

DR WPI: 2000-687541/67.
 DR N-PSDB; AAC81555.
 PT Growth factor homologs and the nucleic acids that encode them, useful
 PT e.g. for treating liver damage, ischemia, multiple sclerosis and
 PT Alzheimer's disease -
 PS Claim 1; Page 110-111; 143pp; English.
 XX
 CC The invention relates to the human growth factor homologue zvegf4
 CC (AAB4653), and nucleic acids encoding it (AAC81555). Zvegf4 is a member
 CC of the PDGF (platelet-derived growth factor)/VEGF (vascular endothelial
 CC growth factor) family. Zvegf4 has a growth factor domain (AAB4654)
 CC characterised by a PDGF cysteine knot structure, and a CUB domain
 CC (AAB4655) which has a beta barrel structure. Zvegf4 has PDGF-like
 CC activity, having mitogenic activity on fibroblasts, vascular smooth
 CC muscle cells and pericytes, and has also been shown to stimulate bone
 CC growth. The invention also relates to fusion proteins comprising human
 CC zvegf4 or fragments thereof, particularly human zvegf4/human zvegf3
 CC fusions; expression constructs and host cells comprising human zvegf4
 CC nucleic acids; the recombinant expression of human zvegf4; an antibody
 CC which binds to human zvegf4 or a fragment thereof; a method of activating
 CC a cell-surface PDGF receptor using a zvegf4-derived polypeptide; a
 CC method of modulating the proliferation, differentiation, migration or
 CC metabolism of bone cells, comprising exposing bone cells to
 CC zvegf4-derived polypeptides; and a method of detecting a genetic
 CC abnormality in the zvegf4 gene of a patient. Zvegf4 proteins and derived
 CC fragments may be used to stimulate tissue development or repair, or
 CC cellular differentiation or proliferation. They are particularly used for
 CC the treatment or repair of liver damage, and may also be used to
 CC modulate neurite growth (e.g., in the treatment of Alzheimer's disease or
 CC multiple sclerosis). Due to their osteogenic activity, they may be used
 CC in the treatment of periodontal disease and fractures. They may be used
 CC to enhance expansion and mobilisation of haematopoietic stem cells
 CC and endothelial precursor stem cells, which may be useful in the
 CC treatment of ischaemia, in wound healing, and in the modulation of the
 CC immune system. The present sequence represents human zvegf4.
 CC
 XX
 SQ Sequence 370 AA;
 Query Match 100.0%; Score 1994; DB 21; Length 370;
 Best Local Similarity 100.0%; Pred. No. 1,1e-188;
 Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MHRLLFYTLICANFCGCRSTATPOSASIKALRNANLRDESHNLTDLRRDETIOYKG 60
 DB 1 MHRLLFYTLICANFCGCRSTATPOSASIKALRNANLRDESHNLTDLRRDETIOYKG 60
 QY 61 NGVQSPRPFNSTPRNLLTWRLHSDENTRIQLVFDNQFLEENNDICRYDFEVEDIS 120
 DB 61 NGVQSPRPFNSTPRNLLTWRLHSDENTRIQLVFDNQFLEENNDICRYDFEVEDIS 120
 QY 121 ETSITIRGRWCGHKEVPPRIKSRTOIKITFKSDDYFAVAPGKRIYSLLEDDQPAASE 180
 DB 121 ETSITIRGRWCGHKEVPPRIKSRTOIKITFKSDDYFAVAPGKRIYSLLEDDQPAASE 180
 QY 181 TNEESTYSSISGVSYNSPSTDTLLADALDKIAEDYEDLLKTFNPESWQDLENNY 240
 DB 181 TNEESTYSSISGVSYNSPSTDTLLADALDKIAEDYEDLLKTFNPESWQDLENNY 240
 QY 241 LDPFRYRGRSYHDKRSKAVDRLNDADAKRYSCTPRNYSVIRBELKIANVYFPRCLLYO 300
 DB 241 LDPFRYRGRSYHDKRSKAVDRLNDADAKRYSCTPRNYSVIRBELKIANVYFPRCLLYO 300
 QY 301 RCGGNGCGGTVMNRSCNCGKTVKKYHEVLAQEPGHIKRGKRAKTMALVDIOLDHHERC 360
 DB 301 RCGGNGCGGTVMNRSCNCGKTVKKYHEVLAQEPGHIKRGKRAKTMALVDIOLDHHERC 360
 QY 361 DCICSSRP 370
 DB 361 DCICSSRP 370

RESULT 2
 ID AAY96864
 XX AAY96864 standard; Protein: 370 AA.
 AC AAY96864;
 XX
 DT 26-SEP-2000 (first entry)
 DE SEQ. ID. 37 from W00034474.
 XX
 KW Vascular endothelial growth factor; homologue; zvegf3; CUB domain;
 KW Cysteine knot; platelet-derived growth factor; PDGF; neuropilin;
 KW chromosome 4q28.3; cytosolic; anti-psoriatic; anti-inflammatory;
 KW anti-diabetic; ophthalmological; anti-rheumatic; anti-arthritis;
 XX
 OS Homo sapiens.
 XX
 FN W0200034474-A2.
 PD 15-JUN-2000.
 XX
 PE 07-DEC-1999; 99WO-0528968.
 XX
 PR 07-DEC-1998; 98US-0207120.
 PR 06-JUL-1999; 99US-0142576.
 PR 21-OCT-1999; 99US-0161653.
 PR 12-NOV-1999; 99US-0165255.
 XX
 PA (ZYMO) ZYMOGENETICS INC.
 XX
 PI Gao Z, Hart CE, Piddington CS, Sheppard PO, Shoemaker KE;
 PI Gilbertson DG, West JW;
 XX
 DR WPI: 2000-423420/36.
 DR N-PSDB; AAA51541.
 XX
 PT Novel zvegf3 polypeptides and nucleotides encoding them useful for
 PT stimulating growth of smooth muscle cells and fibroblasts comprising an
 PT epitope bearing portion of a specific amino acid sequence
 PS Disclosure; Page 164-165; 173pp; English.
 XX
 CC Polypeptides comprising an epitope-bearing portion human or murine
 CC ZVEGF3 (vascular endothelial growth factor homologue) are claimed. The
 CC growth factors comprise a growth factor domain and a CUB domain (generic
 CC sequence motifs are shown in AAY96859 and AAY96860). The growth factor
 CC domain is characterized by an arrangement of cysteine residues and
 CC beta-strands that is characteristic of the "cysteine knot" structure of
 CC the platelet-derived growth factor (PDGF) family. The CUB domain shows
 CC homology to CUB domains in neuropilins, human bone morphogenetic
 CC protein-1, porcine seminal plasma protein, bovine acidic seminal fluid
 CC protein and Xenopus laevis tollold-like protein. Structural analysis and
 CC homology predict that ZVEGF3 polypeptides complex with a second
 CC polypeptide to form multimeric proteins. The human zvegf3 gene has been
 CC mapped to chromosome 4q28.3. ZVEGF3 is useful for stimulating the growth
 CC of fibroblasts or smooth muscle cells, for activating cell surface
 CC PDGF-alpha receptor and for inhibiting PDGF-alpha receptor mediated
 CC cellular processes. ZVEGF3 is useful for regulating (post-development)
 CC organ growth, regeneration and maintenance, as well as tissue
 CC maintenance and repair processes. ZVEGF3 antagonists are useful for
 CC treating cancer, rheumatoid arthritis, diabetic retinopathy, ischemic
 CC limb disease, peripheral vascular disease, myocardial ischemia, vascular
 CC intimal hyperplasia, atherosclerosis, wound healing, chronic liver
 CC disease and haemangioma formation. ZVEGF3 can also be used to modulate
 CC neurite growth and development of the nervous system, and for treating
 CC neurodegenerative diseases.
 XX
 SQ Sequence 370 AA;
 Query Match 100.0%; Score 1994; DB 21; Length 370;
 Best Local Similarity 100.0%; Pred. No. 1,1e-188;
 Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

QY 1 MHRLEFVYTLICANFCSCRDTSATPOSASIKALRNANLRDSENLITDLYRDETIQYKG 60
DB 1 MHRLEFVYTLICANFCSCRDTSATPOSASIKALRNANLRDSENLITDLYRDETIQYKG 60
QY 61 NGYQSPRPNSYPNRLLLTWRLHSQENTRIQLVFNQFGLAEANDICRYDFEVEDIS 120
DB 61 NGYQSPRPNSYPNRLLLTWRLHSQENTRIQLVFNQFGLAEANDICRYDFEVEDIS 120
QY 121 ETSITIRGRCGKHKEVPRIKSRITNOIKITFKSDDYFAKPGFKIYYSLEDFOPAAASE 180
DB 121 ETSITIRGRCGKHKEVPRIKSRITNOIKITFKSDDYFAKPGFKIYYSLEDFOPAAASE 180
QY 181 TNMESVTSISGSVNSPSVDPPTLIADALDKKIAEPTVEDLLKYFPESQEDLENMY 240
DB 181 TNMESVTSISGSVNSPSVDPPTLIADALDKKIAEPTVEDLLKYFPESQEDLENMY 240
QY 241 LDPFRYGRSYHDKRSKYDLDRLNDADAKRYSCPTPNYSVINREELKLANVVEFPCLLVQ 300
DB 241 LDPFRYGRSYHDKRSKYDLDRLNDADAKRYSCPTPNYSVINREELKLANVVEFPCLLVQ 300
QY 301 RCGNCGCGTYVNMRSCTCNSGKTYKKYHEVLQFEPGHIKRRGRATMALVDIQLDHERC 360
DB 301 RCGNCGCGTYVNMRSCTCNSGKTYKKYHEVLQFEPGHIKRRGRATMALVDIQLDHERC 360
QY 361 DCICSSRP 370
DB 361 DCICSSRP 370

```

RESULT 3
AA71130
ID AA71130 standard; Protein: 370 AA.

08-SEP-2000 (first entry)

Human Platelet Derived Growth Factor (PDGF)-D protein.

Platelet Derived Growth Factor-D; PDGF-D; human; cytosolic; vulnary;
VEGF-G; Vascular Endothelial Growth Factor; antiatherosclerotic; tumour;
proliferative; activator; proliferation; differentiation; motility;
growth; PDGF-D receptor; antagonist; tissue remodelling; treat;
atherosclerosis; wound; metastasis.

Homo sapiens.

Key Location/Qualifiers
52..170
/label= CUB domain
/note= "Participates in protein-protein or carbohydrate
interactions"
254..257
/label= Proteolytic site
/note= "Dibasic motif"

Cleavage-site

WO200027879-A1.

18-MAY-2000.

10-NOV-1999; 99WO-US26462.

10-NOV-1998; 98US-0107852.

28-DEC-1998; 98US-0113997.

26-AUG-1999; 99US-0150604.

04-OCT-1999; 99US-0157108.

05-OCT-1999; 99US-0157756.

(LUDM-) LUDWIG INST CANCER RES.
(UHE-) UNIV HELSINKI LICENSING LTD OY.

Eriksson U, Aase K, Fonten A, Lee X, Uutela M, Alltalo K;

Oestman A, Heldin C;
WPI: 2000-376495/32.
N-PSDB: AAD00738.

Novel polynucleotides encoding a novel growth factor of cells
expressing a platelet-derived growth factor, useful for diagnostic and
therapeutic applications, e.g. concerning cancer -

Claim 25; Fig 8; 11pp; English.

The present sequence is the complete human platelet derived growth factor
(PDGF)-D, formally known as Vascular Endothelial Growth Factor (VEGF)-G.
It is derived from human foetal lung lamdaglio cDNA library. It belongs
to the VEGF/PDGF family. It functions as an activator of proliferation,
differentiation, growth and motility of cells, that express PDGF-D
receptor. This sequence is useful for inhibiting the growth of tumours,
that express PDGF-D. Expression of PDGF-D and its proteolytic cleavage
for generating an activated truncated form is useful for regulating
receptor binding specificity of PDGF-D. PDGF-D antagonist is useful for
inhibiting tissue remodelling during the invasion of tumour cells into
normal cells. PDGF-D may be used to treat wounds, atherosclerosis,
metastasis and migration of smooth muscle cells.

Sequence 370 AA:

Query Match 100.0%; Score 1994; DB 21; Length 370;
Best Local Similarity 100.0%; Pred. No. 1.le-188;
Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

QY 1 MHRLEFVYTLICANFCSCRDTSATPOSASIKALRNANLRDSENLITDLYRDETIQYKG 60
DB 1 MHRLEFVYTLICANFCSCRDTSATPOSASIKALRNANLRDSENLITDLYRDETIQYKG 60
QY 61 NGYQSPRPNSYPNRLLLTWRLHSQENTRIQLVFNQFGLAEANDICRYDFEVEDIS 120
DB 61 NGYQSPRPNSYPNRLLLTWRLHSQENTRIQLVFNQFGLAEANDICRYDFEVEDIS 120
QY 121 ETSITIRGRCGKHKEVPRIKSRITNOIKITFKSDDYFAKPGFKIYYSLEDFOPAAASE 180
DB 121 ETSITIRGRCGKHKEVPRIKSRITNOIKITFKSDDYFAKPGFKIYYSLEDFOPAAASE 180
QY 181 TNMESVTSISGSVNSPSVDPPTLIADALDKKIAEPTVEDLLKYFPESQEDLENMY 240
DB 181 TNMESVTSISGSVNSPSVDPPTLIADALDKKIAEPTVEDLLKYFPESQEDLENMY 240
QY 241 LDPFRYGRSYHDKRSKYDLDRLNDADAKRYSCPTPNYSVINREELKLANVVEFPCLLVQ 300
DB 241 LDPFRYGRSYHDKRSKYDLDRLNDADAKRYSCPTPNYSVINREELKLANVVEFPCLLVQ 300
QY 301 RCGNCGCGTYVNMRSCTCNSGKTYKKYHEVLQFEPGHIKRRGRATMALVDIQLDHERC 360
DB 301 RCGNCGCGTYVNMRSCTCNSGKTYKKYHEVLQFEPGHIKRRGRATMALVDIQLDHERC 360
QY 361 DCICSSRP 370
DB 361 DCICSSRP 370

```

RESULT 4
AAG65601
ID AAG65601 standard; Protein: 370 AA.

AAG65601;

07-JAN-2002 (first entry)

Human zveg4 polypeptide.

zveg4; bone; ligament; cartilage; osteoblast; osteoclast; chondrocyte;
bone cancer; osteonecrosis; bone defect; osteogenesis; osteoporosis;
osteopathic; vulnary; human.

OS Homo sapiens.
 XX Key Location/Qualifiers
 FT Peptide 1..18
 FT Protein /note= "secretory peptide"
 FT Protein 19..370
 FT Domain /note= "mature protein"
 FT Domain 52..179
 FT Region /note= "CUB domain"
 FT Region 180..257
 FT Domain /note= "propeptide-like sequence"
 FT Domain 258..370
 FT Domain /note= "growth factor domain"
 PN WO200157083-A1.
 XX 09-AUG-2001.
 XX 03-MAY-2000; 2000WO-US12095.
 XX 04-FEB-2000; 2000US-180169P.
 PR 31-MAR-2000; 2000US-0540224.
 XX (ZYMO) ZYMOGENETICS INC.
 XX Gilbertson DG, Hart CE;
 PI WPI; 2001-611088/70.
 DR N-PSDB; AAA47772.
 XX
 PT Use of zvegf4 polypeptide for promoting bone, ligament or cartilage
 PT growth in mammal at site of fracture, implant, and bone graft, and for
 PT promoting growth or differentiation of osteoblasts, chondrocytes in
 PT culture
 XX
 PS Example 2; Page 44-47; 57pp; English.
 XX
 CC The invention relates to the use of zvegf4 polypeptide for promoting
 CC bone, ligament or cartilage growth in a mammal, and for promoting
 CC proliferation or differentiation of osteoblasts, osteoclasts,
 CC chondrocytes or bone marrow stem cells in culture. For promoting
 CC cartilage growth, chondrocytes are cultured ex vivo in presence of the
 CC zvegf4 polypeptide and then placed into mammal where cartilage is to be
 CC grown. Zvegf4 polypeptide is useful for promoting growth of bone,
 CC ligament or cartilage in a mammal at a site of bony defect such as
 CC fracture, bone graft, implant or periodontal pocket, in humans and non-
 CC human animals such as domestic animals including livestock and companion
 CC animals. Zvegf4 is used for promoting growth of bone, ligament, or
 CC cartilage in conditions of bone defects following therapeutic treatments
 CC of bone cancers or other conditions characterized by increased bone loss
 CC or decreased bone formation, or elevation of peak bone mass in pre-
 CC menopausal women. It is also useful for healing bone following radiation
 CC -induced osteonecrosis, repairing bone defects arising from surgery, and
 CC promotion of bone healing in plastic surgery, increasing bone formation
 CC during distraction osteogenesis, treating bone injuries including repair
 CC of cartilage and ligament and treatment of osteoporosis. The present
 CC sequence represents a human zvegf4 polypeptide.
 CC
 CC Sequence 370 AA:
 SQ
 Query Match 100.0%; Score 1994; DB 22; Length 370;
 Best Local Similarity 100.0%; Pred. No. 1,1e-188;
 Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MHRLLFVYLICANFCSCRDTSATPOSASIKALRNANLRDSENNHLDYRRRETOVG 60
 DB 1 MHRLLFVYLICANFCSCRDTSATPOSASIKALRNANLRDSENNHLDYRRRETOVG 60
 QY 61 NCYVSPRPNSYPRLNLTWRLHSGENRIQLVDPNQGFLAEANDICRYFVEVEDS 120
 DB 61 NCYVSPRPNSYPRLNLTWRLHSGENRIQLVDPNQGFLAEANDICRYFVEVEDS 120
 QY 121 ESTTIRGRMGCHKEVPPRIKSRITNOIKTFKSDDYFAKPGFKIYSLLEDPQPAASE 180
 DB 121 ESTTIRGRMGCHKEVPPRIKSRITNOIKTFKSDDYFAKPGFKIYSLLEDPQPAASE 180

DB 121 ESTTIRGRMGCHKEVPPRIKSRITNOIKTFKSDDYFAKPGFKIYSLLEDPQPAASE 180
 QY 181 TMNESVTSSISGYSNPSVTDPLLDALDKTAEPDVEDDLKYNPESWQEDLEMY 240
 DB 181 TMNESVTSSISGYSNPSVTDPLLDALDKTAEPDVEDDLKYNPESWQEDLEMY 240
 QY 241 LDTPRYGRSYHDKRSKYVDLDRNLNDAKRYSCPTPNTSVNIREELKLANVYFPRCLLYQ 300
 DB 241 LDTPRYGRSYHDKRSKYVDLDRNLNDAKRYSCPTPNTSVNIREELKLANVYFPRCLLYQ 300
 QY 301 RCGNCGCGTYNMRSCTNSGKTYKKYHEVLOEPEPGIKRGRAKTALVDIQLDHERC 360
 DB 301 RCGNCGCGTYNMRSCTNSGKTYKKYHEVLOEPEPGIKRGRAKTALVDIQLDHERC 360
 QY 361 DCICSSRPPR 370
 DB 361 DCICSSRPPR 370
 RESULT 5
 AAB85529 standard; protein; 370 AA.
 AAB85529;
 25-SEP-2001 (first entry)
 Human secreted protein (clone id HGCNC48).
 Secreted protein; immunosuppressive; antiarthritic; antirheumatic;
 antiapoptotic; cytosolic; cardiant; vasotropic; cerebroprotective;
 neurotropic; neuroprotective; antibacterial; virucide; fungicide; human;
 ophthalmological; gene therapy.
 Homo sapiens.
 WO200155430-A1.
 02-AUG-2001.
 17-JAN-2001; 2001WO-US01431.
 31-JAN-2000; 2000US-0179065.
 04-FEB-2000; 2000US-0180628.
 12-SEP-2000; 2000US-0231968.
 (HUMA-) HUMAN GENOME SCI INC.
 Rosen CA, Komatsu J, Baker KP, Birse CE, Soppet DR, Olsen HS;
 Moore PA, Wei P, Ebner R, Duan DR, Shi Y, Choi GH, Fiscella M;
 Ni J, Ruben SM, Barash SC;
 WPI; 2001-476220/51.
 N-PSDB; AAA46939.
 17 isolated nucleic acid molecules encoding human secreted proteins,
 used to preventing, treating or ameliorating a medical condition
 Claim 11; Page 447-449; 482pp; English.
 The invention provides novel human secreted proteins and polynucleotides
 encoding them. The secreted proteins can be expressed by standard
 recombinant methodology. The secreted proteins and polynucleotides are
 used to prevent, treat or ameliorate a medical condition in e.g. humans,
 mice, rabbits, goats, horses, cats, dogs, chickens or sheep. They can
 also be used in diagnosing a pathological condition. The antibodies to
 the proteins can also be used in alleviating symptoms associated with the
 disorders and in diagnostic immunoassays e.g. radioimmunoassays or enzyme
 linked immunosorbent assays (ELISA). Disorders which are diagnosed or
 treated include autoimmune diseases e.g. Rheumatoid arthritis,
 hyperproliferative disorders e.g. neoplasms of the breast or liver,
 cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders

CC e.g. cerebral ischemia, angiogenesis, nervous system disorders e.g.
 CC Alzheimer's disease, infections caused by bacteria, viruses and fungi and
 CC ocular disorders e.g. corneal infection. The polypeptides can also be
 CC used to aid wound healing and epithelial cell proliferation, to prevent
 CC skin aging due to sunburn, to maintain organs before transplantation, for
 CC supporting cell culture of primary tissues, to regenerate tissues and in
 CC chemotaxis. The polypeptides can also be used as a food additive or
 CC preservative to increase or decrease storage capabilities. The present
 CC sequence represents a human secreted protein.

XX Sequence 370 AA:

Query Match 100.0%; Score 1994; DB 22; Length 370;
 Best Local Similarity 100.0%; Pred. No. 1,1e-188;
 Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MHRLLFYVTLICANFCSCRDTSATPQASIKALRNANLRDSENLTLDRDETIOVKG 60
 DB 1 MHRLLFYVTLICANFCSCRDTSATPQASIKALRNANLRDSENLTLDRDETIOVKG 60
 QY 61 NGYVSPRPNSYPNNLLTWRLHSQENTRIQLVFDNPGLEAENDICRYDFVEYEDIS 120
 DB 61 NGYVSPRPNSYPNNLLTWRLHSQENTRIQLVFDNPGLEAENDICRYDFVEYEDIS 120
 QY 121 ESTTIRGRMGCKHKEVPPRIKSRNTQIKTFKSDYFYAKPGFKIYSLLEDFOPAAASE 180
 DB 121 ESTTIRGRMGCKHKEVPPRIKSRNTQIKTFKSDYFYAKPGFKIYSLLEDFOPAAASE 180
 QY 181 TNMESVTSISGSVSNPSVTDPTLLADALDKKIAEPTVEDLLKYFNDESQEDLENNY 240
 DB 181 TNMESVTSISGSVSNPSVTDPTLLADALDKKIAEPTVEDLLKYFNDESQEDLENNY 240
 QY 241 LDTPYRGRSHYDRKSKVDRLNDARKRYCTPRNYSYNIEEKLAVNVPFPCLLVQ 300
 DB 241 LDTPYRGRSHYDRKSKVDRLNDARKRYCTPRNYSYNIEEKLAVNVPFPCLLVQ 300
 QY 301 RCGNGCGGTVMNRSCCTNSGKTVKKYHEVLQFEFGHIIKRRGRATMALVDIQLDHHERC 360
 DB 301 RCGNGCGGTVMNRSCCTNSGKTVKKYHEVLQFEFGHIIKRRGRATMALVDIQLDHHERC 360
 QY 361 DCICSSRPPR 370
 DB 361 DCICSSRPPR 370

RESULT 6

AAU00698 standard; Protein: 370 AA.

AC AAU00698;

DT 07-SEP-2001 (first entry)

DE Human FCTR1 protein present in clone 30664188.0.99.

XX Bone morphogenetic protein-1; BMP-1; Vascular endothelial growth factor;
 KW VEGF-E; platelet derived growth factor; PDGF; FCTR1; hyperplasia; cancer;
 KW neoplasia; anaemia; leukopenia; baldness; cardiovascular disorder;
 KW fibrotic disorder; diabetic ulcer; obesity; hyperproliferation; human;
 KW dysproliferation; neurodegenerative disorder; osteoarthritis; epilepsy;
 KW inflammatory disorder; Graft versus host disease; coagulation;
 KW haemophilia; neural disorder; Parkinson's disease; Alzheimer's disease;
 KW multiple sclerosis; Huntington's disease; amyotrophic lateral sclerosis;
 KW peripheral neuropathy; acute brain injury.

XX Homo sapiens.

OS Key Location/Qualifiers

FT Peptide 1..23

FT Protein /note= "Signal peptide"

FT /note= "Mature FCTR1"

FT Domain 53..167

FT /note= "CUB domain"
 FT Domain 272..306
 FT /note= "PDGF domain"
 FT Modified-site 276
 FT /note= "N-linked glycosylation site"
 FT Domain 302..365
 FT /note= "Metallothionein domain"
 FT Domain 350..362
 FT /note= "PDGF domain"

PN WO200125437-A2.

XX 12-APR-2001.

XX 06-OCT-2000; 2000WO-US27671.

XX 07-OCT-1999; 99US-0158083.

XX 13-OCT-1999; 99US-0159231.

XX 04-JAN-2000; 2000US-0174485.

XX 03-MAR-2000; 2000US-0186707.

XX 10-MAR-2000; 2000US-0188250.

XX 08-AUG-2000; 2000US-0223879.

XX 12-SEP-2000; 2000US-0662783.

XX 20-SEP-2000; 2000US-0234082.

XX (CURA-) CURAGEN CORP.

XX Shinketsu RA, Lichenstein H, Herrmann JL, Boldog FL, Minskoff S;
 PI Jeffers M;

XX WPI: 2001-316172/73.

XX N-PSDB: AAS04492.

XX Novel growth factor polypeptides termed as FCTR1 polypeptides, useful
 PT for treating cancer, cardiovascular and fibrotic diseases, diabetic
 PT ulcers, wound healing and neuronal disorders

PS Claim 1; Fig 1; 17pp; English.

CC The sequence represents a protein related to bone morphogenetic protein-1
 CC (BMP-1), vascular endothelial growth factor (VEGF-E) and platelet derived
 CC growth factor (PDGF). Polypeptides and polynucleotides related to BMP-1,
 CC VEGF-E and PDGF are referred to as FCTR1 peptides and nucleic acids.
 CC FCTR1 proteins are useful for treating or preventing a disorder
 CC associated with aberrant expression, aberrant processing, or aberrant
 CC physiological interactions of the proteins in a mammal, where the
 CC disorder is characterised by insufficient or ineffective growth of a cell
 CC or a tissue, e.g. hyperplasia or neoplasia. The peptides and their
 CC associated nucleic acids are useful for both promoting and inhibiting
 CC growth of cells and tissues and in treatment of cancer, anaemia,
 CC leukopenia, baldness, for treating cardiovascular and fibrotic disorders,
 CC diabetic ulcers, obesity, infectious diseases, hyperproliferative and
 CC dysproliferative disorders, neurodegenerative disorders, osteoarthritis,
 CC inflammatory disorders, Graft versus host disease, coagulation disorders
 CC such as haemophilia, and neural disorders including Parkinson's disease,
 CC Alzheimer's disease, multiple sclerosis, Huntington's disease,
 CC amyotrophic lateral sclerosis, peripheral neuropathy, acute brain injury
 CC and epilepsy.

XX Sequence. 370 AA:

QY Query Match 100.0%; Score 1994; DB 22; Length 370;

DB Best Local Similarity 100.0%; Pred. No. 1,1e-188;
 Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MHRLLFYVTLICANFCSCRDTSATPQASIKALRNANLRDSENLTLDRDETIOVKG 60

DB 1 MHRLLFYVTLICANFCSCRDTSATPQASIKALRNANLRDSENLTLDRDETIOVKG 60

QY 61 NGYVSPRPNSYPNNLLTWRLHSQENTRIQLVFDNPGLEAENDICRYDFVEYEDIS 120

DB 61 NGYVSPRPNSYPNNLLTWRLHSQENTRIQLVFDNPGLEAENDICRYDFVEYEDIS 120


```

Db 61 NGVOSPPFPNSYPNNLLITWRLHSQENTRIQLVFDNPGLEAENDICRYDFVEEDIS 120
QY 121 ETSIIIRGRMCHEKPEVPRKSRITNOIKITFKSDYFVAKPGFKIYSLLEDFOPAAASE 180
Db 121 ETSIIIRGRMCHEKPEVPRKSRITNOIKITFKSDYFVAKPGFKIYSLLEDFOPAAASE 180
QY 181 TNWESVTSISGVSNSPSYDPTLIADALDKKIAEFTVEEDLYKFNESQOEDLENNY 240
Db 181 TNWESVTSISGVSNSPSYDPTLIADALDKKIAEFTVEEDLYKFNESQOEDLENNY 240
QY 241 LDTPRYGRGRSYHDKRSKYDLRLNDADAKRYSCTPRANYSVINIREELKLANVFFPCLLYQ 300
Db 241 LDTPRYGRGRSYHDKRSKYDLRLNDADAKRYSCTPRANYSVINIREELKLANVFFPCLLYQ 300
QY 301 RCGNGCGGTYNWSCTNSGKTYKKYHEVLQFEPGHIKRRGRAKTMALVLDIQLDHHERC 360
Db 301 RCGNGCGGTYNWSCTNSGKTYKKYHEVLQFEPGHIKRRGRAKTMALVLDIQLDHHERC 360
QY 361 DCICSSRPPR 370
Db 361 DCICSSRPPR 370

RESULT 10
ABP51640
ID ABP51640 standard; Protein: 370 AA.
AC ABP51640;
XX 30-SEP-2002 (first entry)
XX Human zvegfg4 protein SEQ ID NO:2.
DE
XX
XX Human; zvegfg4; cell proliferation; extracellular matrix production;
XX fibroproliferative disorder; PDGF-D; platelet derived growth factor;
XX PDGF; vascular endothelial growth factor; VEGF; cytostatic; nephrotropic;
XX hepatotropic; antiinflammatory; osteopathic; antiarthritic; metastasis;
XX prostate tumor; prostate cancer; glomerulonephritis; lupus nephritis;
XX diabetic glomerulosclerosis; renal arteriosclerosis; nephrotic syndrome;
XX chronic active hepatitis; cirrhosis; osteopetrosis; osteosclerosis;
XX hyperostosis; osteoarthritis.
XX
XX Homo sapiens.
XX
XX OS
XX PN US2002064832-A1.
XX
XX PD 30-MAY-2002.
XX
XX PF 14-MAR-2001; 2001US-0808972.
XX
XX PR 03-MAY-1999; 99US-132250P.
XX PR 10-NOV-1999; 99US-164463P.
XX PR 04-FEB-2000; 2000US-180169P.
XX PR 26-SEP-2000; 2000US-235295P.
XX PR 03-MAY-2000; 2000US-0564595.
XX
XX PA (HART/) HART C E.
XX PA (TOPO/) TOPOUZIS S.
XX PA (GILB/) GILBERTSON D G.
XX
XX PI Hart CE, Topouzis S, Gilbertson DG;
XX
XX DR WPI: 2002-573696/61.
XX DR N-PSDB; ABQ73239.
XX
XX PT Reducing proliferation or extracellular matrix production by a cell in
XX liver and kidney, comprises administering a zvegfg4 antagonist
XX
XX PS Example 3; Page 19-20; 34pp; English.
XX
XX CC The present invention describes a method for reducing proliferation of

```

```

CC or extracellular matrix production by a cell in a mammal. The method
CC comprises administering to the mammal a composition comprising a
CC therapeutically effective amount of a zvegfg4 antagonist chosen from
CC anti-zvegfg4 antibodies, inhibitory polynucleotides, inhibitors of
CC zvegfg4 activation, and mitogenically inactive, receptor-binding variants
CC of zvegfg4. Zvegfg4 (also called PDGF-D) is a multi-domain protein that is
CC structurally related to platelet derived growth factor (PDGF) and
CC vascular endothelial growth factors (VEGF). Zvegfg4 has cytostatic,
CC nephrotropic, hepatotropic, antiinflammatory, osteopathic and
CC antiarthritic activities. The method is useful for reducing proliferation
CC of mesangial, epithelial, endothelial, smooth muscle, fibroblast,
CC osteoblast, osteoclast, neuronal, stromal, stellate or interstitial cells
CC in a mammal, in particular proliferation of prostate tumour cells, and
CC for reducing extracellular matrix production by a cell in a mammal
CC suffering from a fibroproliferative disorder of kidney, bone or liver.
CC In particular it is useful for reducing stellate cell activation. The
CC method is useful for reducing metastasis of prostate cancer cells to
CC bone in a mammal and for treating a fibroproliferative disorder of
CC kidney, liver or bone in a mammal. Fibroproliferative disorders of the
CC kidney include, glomerulonephritis, diabetic glomerulosclerosis, lupus
CC nephritis, renal arteriosclerosis and nephrotic syndrome, disorders of
CC the liver include chronic active hepatitis and many other types of
CC cirrhosis, and disorders of the bone include osteopetrosis, hyperostosis,
CC osteosclerosis, osteoarthritis, and ectopic bone formation in metastatic
CC prostate cancer. The present sequence represents human zvegfg4, which is
CC used in an example from the present invention.
XX
XX SQ Sequence 370 AA;

```

```

Query Match 100.0%; Score 1994; DB 23; Length 370;
Best Local Similarity 100.0%; Pred. No. 1,1e-188;
Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

QY 1 MHRLLFYVTLICANFSCSDTSATFQASISIKALRNANLRDPSNHLTLYRDETIQYKG 60
Db 1 MHRLLFYVTLICANFSCSDTSATFQASISIKALRNANLRDPSNHLTLYRDETIQYKG 60
QY 61 NGVOSPPFPNSYPNNLLITWRLHSQENTRIQLVFDNPGLEAENDICRYDFVEEDIS 120
Db 61 NGVOSPPFPNSYPNNLLITWRLHSQENTRIQLVFDNPGLEAENDICRYDFVEEDIS 120
QY 121 ETSIIIRGRMCHEKPEVPRKSRITNOIKITFKSDYFVAKPGFKIYSLLEDFOPAAASE 180
Db 121 ETSIIIRGRMCHEKPEVPRKSRITNOIKITFKSDYFVAKPGFKIYSLLEDFOPAAASE 180
QY 181 TNWESVTSISGVSNSPSYDPTLIADALDKKIAEFTVEEDLYKFNESQOEDLENNY 240
Db 181 TNWESVTSISGVSNSPSYDPTLIADALDKKIAEFTVEEDLYKFNESQOEDLENNY 240
QY 241 LDTPRYGRGRSYHDKRSKYDLRLNDADAKRYSCTPRANYSVINIREELKLANVFFPCLLYQ 300
Db 241 LDTPRYGRGRSYHDKRSKYDLRLNDADAKRYSCTPRANYSVINIREELKLANVFFPCLLYQ 300
QY 301 RCGNGCGGTYNWSCTNSGKTYKKYHEVLQFEPGHIKRRGRAKTMALVLDIQLDHHERC 360
Db 301 RCGNGCGGTYNWSCTNSGKTYKKYHEVLQFEPGHIKRRGRAKTMALVLDIQLDHHERC 360
QY 361 DCICSSRPPR 370
Db 361 DCICSSRPPR 370

```

```

RESULT 11
ABG64733
ID ABG64733 standard; Protein: 370 AA.
XX
XX AC ABG64733;
XX
XX DT 27-AUG-2002 (first entry)
XX
XX DE Human albumin fusion protein #1408.
XX
XX KW Albumin fusion protein; therapeutic protein X; human albumin; HA;

```


KM human serum albumin; HSA; cancer; reproductive disorder;
 KM digestive disorder; immune disorder; endocrine disorder;
 KM haematopoietic disorder; neural disorder; connective disorder;
 KM cytostatic; antineoplastic; antiinflammatory; anticancer;
 KM immunomodulator; anti-HIV; antidiabetic; haemostatic; nootropic;
 KM neuroprotective; antiparkinsonian; antimicrobial; neuroleptic;
 KM osteopathic; antiarthritic.
 XX
 OS Homo sapiens.
 OS Synthetic.
 XX
 PN WO200177137-A1.
 XX
 PD 18-OCT-2001.
 XX
 PF 12-APR-2001; 2001WO-US11988.
 XX
 PR 12-APR-2000; 2000US-229358P.
 XX
 PR 25-APR-2000; 2000US-199384P.
 XX
 PR 21-DEC-2000; 2000US-256931P.
 XX
 PA (HUMA-) HUMAN GENOME SCI INC.
 XX
 PI Rosen CA, Haseltine WA;
 XX
 DR WPI: 2002-010886/01.
 XX
 PT New fusion protein for treating disease e.g. diabetes comprises an
 PT albumin fused to a therapeutic protein -
 XX
 PS Claim 1; Page 1459-1460; 2102pp; English.
 XX
 CC The present invention relates to albumin fusion proteins comprising a
 CC therapeutic protein X and human albumin (HA, also known as human serum
 CC albumin, HSA). The proteins are useful for treating a disease or
 CC disorder that may be modulated by therapeutic protein X. The albumin
 CC extends the shelf-life of protein X, and may increase its biological
 CC in vitro/in vivo activity. The protein is useful for treating and
 CC diagnosing disorders such as cancer, reproductive disorders, digestive
 CC disorders (e.g. Crohn's disease, ulcerative colitis), immune disorders
 CC (e.g. acquired immunodeficiency syndrome, AIDS), endocrine disorders
 CC (e.g. diabetes), haematopoietic disorders, neural disorders
 CC (e.g. Alzheimer's, Parkinson's, Creutzfeldt-Jacob disease,
 CC encephalomyelitis, meningitis, schizophrenia), and connective disorders
 CC (e.g. osteoporosis, arthritis). ABG63326-ABG63518 represent albumin
 CC fusion proteins of the invention.
 CC
 XX
 SQ Sequence 370 AA:
 Query Match 100.0%; Score 1994; DB 23; Length 370;
 Best Local Similarity 100.0%; Pred. No. 1,1e-188;
 Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MHRLFFVYLICANFCSCRDTSATPOSASIKALRNANLRDESNHLLTDYRDETIQVKG 60
 DB 1 MHRLFFVYLICANFCSCRDTSATPOSASIKALRNANLRDESNHLLTDYRDETIQVKG 60
 QY 61 NGYVSPRPNSYPNNLLTWRLHSOENTRIQVFNQGLEAENDICRYFVEVEDIS 120
 DB 61 NGYVSPRPNSYPNNLLTWRLHSOENTRIQVFNQGLEAENDICRYFVEVEDIS 120
 QY 121 EFTSTIRGRMGCHKEVPRIKSRTOIKTEKSDSYFAVAKPEFKIYISLLEDFOPAAASE 180
 DB 121 EFTSTIRGRMGCHKEVPRIKSRTOIKTEKSDSYFAVAKPEFKIYISLLEDFOPAAASE 180
 QY 181 TMWESVTSISGVSNPSVPTDPTLADALDKKIAEFTVEEDLLKYFNESWQEDLENNY 240
 DB 181 TMWESVTSISGVSNPSVPTDPTLADALDKKIAEFTVEEDLLKYFNESWQEDLENNY 240
 QY 241 LDTPTIRGRSYHDKRSKYDLRLNDADAKYSCTPRNYSVINREELKLANVVEFPRLCLVQ 300
 DB 241 LDTPTIRGRSYHDKRSKYDLRLNDADAKYSCTPRNYSVINREELKLANVVEFPRLCLVQ 300

QY 301 RCGNCGGVTVMWRSCCTNSGKTYKHYEVLQFEPGHKIRGRATMALVDIQLDHHERC 360
 DB 301 RCGNCGGVTVMWRSCCTNSGKTYKHYEVLQFEPGHKIRGRATMALVDIQLDHHERC 360
 QY 361 DCICSSRPPR 370
 DB 361 DCICSSRPPR 370

RESULT 12

AAB47891

ID AAB47891 standard; Protein; 370 AA.

XX AAB47891;

XX 16-MAY-2002 (first entry)

XX Human zvegfg.

XX Human; mouse; zvegfg; platelet derived growth factor;

XX PDGF; homolog; growth; bone; ligament; cartilage; proliferation;

XX implant; periodontal pocket; osteoclast; bone marrow stem cell;

XX osteoporosis.

XX Homo sapiens.

XX US2002004225-A1.

XX 10-JAN-2002.

XX 29-MAR-2001; 2001US-0823033.

XX 07-DEC-1998; 980S-111173P.

XX 06-JUL-1999; 990S-142576P.

XX 21-OCT-1999; 990S-161653P.

XX 12-NOV-1999; 990S-165255P.

XX 31-MAR-2000; 2000US-193723P.

XX 07-DEC-1999; 990S-0457066.

XX (HART/) HART C E.

XX (GILB/) GILBERTSON D G.

XX Hart CE, Gilbertson DG;

XX WPI: 2002-171026/22.

XX Promoting growth of bone, ligament or cartilage in a mammal, involves

XX administering to the mammal a protein which comprises growth factor

XX domain of zvegfg protein, a homolog of platelet-derived growth factor

XX Claim 8; Page 20-21; 31pp; English.

XX This sequences represents human zvegfg. zvegfg can be used in a

XX composition with either human or mouse zvegfg, for promoting growth of

XX bone, ligament or cartilage and stimulating proliferation of osteoblasts

XX or chondrocytes in a mammal. zvegfg is a platelet derived growth factor

XX (PDGF) homolog. The zvegfg protein used was preferably a dimeric

XX protein, with a delivery vehicle. The method of th invention is useful

XX for promoting growth of bone, ligament or cartilage in a mammal, where

XX the composition is administered at a site of a bony defect, preferably

XX a fracture, bone graft site, implant site, or periodontal pocket, and

XX for stimulating proliferation of osteoblasts or chondrocytes in a

XX mammal. It is further useful for promoting proliferation of osteoblasts,

XX osteoclasts, chondrocytes or bone marrow stem cells, where the bone

XX marrow stem cells are harvested from a patient prior to culture. The

XX method is therefore useful for treating osteoporosis.

XX
 XX Sequence 370 AA:
 Query Match 100.0%; Score 1994; DB 23; Length 370;

Best Local Similarity 100.0%; Pred. No. 1,1e-188;
Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

QY 1 MHRLEFVYLLICANFCSCRDTSATPQASISIKALRNANLRDESNHLDLYRDEDTIOYKG 60
Db 1 MHRLEFVYLLICANFCSCRDTSATPQASISIKALRNANLRDESNHLDLYRDEDTIOYKG 60
QY 61 NGYQSPRPNSYPNNLLTWRLHSENTRIQLVFDNFGLEAENDICRYDFEVEDIS 120
Db 61 NGYQSPRPNSYPNNLLTWRLHSENTRIQLVFDNFGLEAENDICRYDFEVEDIS 120
QY 121 ESTIIRGRMCHEVPPRIKSRNQITFKSDDYFAKGGFKIYSLDEFOPAASE 180
Db 121 ESTIIRGRMCHEVPPRIKSRNQITFKSDDYFAKGGFKIYSLDEFOPAASE 180
QY 181 TMSVTSISGISVSNPSVTDPTLIADALDKIAEFTVDLKYFNPESMOEDLEMY 240
Db 181 TMSVTSISGISVSNPSVTDPTLIADALDKIAEFTVDLKYFNPESMOEDLEMY 240
QY 241 LDTPRYGRSYHDKRSKYDLDRLNDADAKRYSCPTRNYSVNIREELKLANVVEFPRLCYQ 300
Db 241 LDTPRYGRSYHDKRSKYDLDRLNDADAKRYSCPTRNYSVNIREELKLANVVEFPRLCYQ 300
QY 301 RCGNCGCGTYNMRSCTCNSGKTYKKYHEVLQFEPGHKRRGRAKTALVDIQLDHHERC 360
Db 301 RCGNCGCGTYNMRSCTCNSGKTYKKYHEVLQFEPGHKRRGRAKTALVDIQLDHHERC 360
QY 361 DCICSSRPPR 370
Db 361 DCICSSRPPR 370

```

RESULT 13

AAE15819
ID AAE15819 standard; Protein; 370 AA.

XX AAE15819;

DT 26-MAR-2002 (first entry)

XX Human LP85 protein #1.

XX LP85; platelet-derived growth factor; PDGF; antiinflammatory; vulnery;
XX osteopathic; neuroprotective; tranquilliser; musculoskeletal disorder;
XX MSD; therapy; bone growth; cartilage differentiation; wound healing;
XX neuron growth; bone fracture; osteoporosis; osteopenia; arthritis;
XX sarcopenia; periodontal disease; tissue atrophy; endocrine disorder;
XX muscle loss; immobility; bone density.

OS Homo sapiens.

XX Homo sapiens.

XX Key Location/Qualifiers

XX Peptide 1..12

XX Protein /label= signal_peptide

XX /label= Human_mature_LP85_protein

XX W0200189450-A2.

XX 29-NOV-2001.

XX 08-MAY-2001; 2001WO-US11755.

XX 19-MAY-2000; 2000US-2054424P.

XX 11-JAN-2001; 2001US-261071P.

XX 11-JAN-2001; 2001US-261076P.

XX (ELIL) LILLY & CO ELI.

XX Beals JM, Gonzales-Dewhitt PA, Hammond LJ, Lu J, Na S, Su EW;

XX Wlitcher DR, Wroblewski VJ;

XX WPI; 2002-083040/L1.

DR N-PSDB; AAE15819.

XX Analog of a platelet-derived growth factor homolog, LP85 useful for
PT treating osteoporosis, arthritis, sarcopenia, wounds, has one or more
PT amino acid substitutions which destroy the tripeptidyl sequence of
PT native LP85

PS Claim 11; Page 109-110; 117pp; English.

CC The present invention relates to LP85, an analogue of platelet-derived
CC growth factor (PDGF) homologue. Sequences of the invention are useful
CC for the manufacture of a medicament for treating musculoskeletal disorder
CC (MSD) which include promoting bone growth, cartilage differentiation and
CC function, wound healing, neuron growth, preventing cartilage degradation
CC or neuronal degeneration. They are useful for treating bone fractures,
CC osteoporosis, osteopenia, arthritis, sarcopenia, periodontal disease,
CC tissue atrophy, traumatised connective tissues, grafted connective
CC tissues and/or transplanted organs, bone or muscle loss due to
CC malignancy, endocrine disorders and immobility. They are also used
CC for prophylactically increasing or maintaining bone density in a
CC mammal. The present sequence is human LP85 protein.

SO Sequence 370 AA;

Query Match 100.0%; Score 194; DB 23; Length 370;
Best Local Similarity 100.0%; Pred. No. 1,1e-188;
Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

QY 1 MHRLEFVYLLICANFCSCRDTSATPQASISIKALRNANLRDESNHLDLYRDEDTIOYKG 60
Db 1 MHRLEFVYLLICANFCSCRDTSATPQASISIKALRNANLRDESNHLDLYRDEDTIOYKG 60
QY 61 NGYQSPRPNSYPNNLLTWRLHSENTRIQLVFDNFGLEAENDICRYDFEVEDIS 120
Db 61 NGYQSPRPNSYPNNLLTWRLHSENTRIQLVFDNFGLEAENDICRYDFEVEDIS 120
QY 121 ESTIIRGRMCHEVPPRIKSRNQITFKSDDYFAKGGFKIYSLDEFOPAASE 180
Db 121 ESTIIRGRMCHEVPPRIKSRNQITFKSDDYFAKGGFKIYSLDEFOPAASE 180
QY 181 TMSVTSISGISVSNPSVTDPTLIADALDKIAEFTVDLKYFNPESMOEDLEMY 240
Db 181 TMSVTSISGISVSNPSVTDPTLIADALDKIAEFTVDLKYFNPESMOEDLEMY 240
QY 241 LDTPRYGRSYHDKRSKYDLDRLNDADAKRYSCPTRNYSVNIREELKLANVVEFPRLCYQ 300
Db 241 LDTPRYGRSYHDKRSKYDLDRLNDADAKRYSCPTRNYSVNIREELKLANVVEFPRLCYQ 300
QY 301 RCGNCGCGTYNMRSCTCNSGKTYKKYHEVLQFEPGHKRRGRAKTALVDIQLDHHERC 360
Db 301 RCGNCGCGTYNMRSCTCNSGKTYKKYHEVLQFEPGHKRRGRAKTALVDIQLDHHERC 360
QY 361 DCICSSRPPR 370
Db 361 DCICSSRPPR 370

```

RESULT 14

AAE15845
ID AAE15845 standard; Protein; 370 AA.

XX AAE15845;

DT 26-MAR-2002 (first entry)

XX Human LP85 mutant protein (N276D).

XX LP85; platelet-derived growth factor; PDGF; antiinflammatory; vulnery;
XX osteopathic; neuroprotective; tranquilliser; musculoskeletal disorder;
XX MSD; therapy; bone growth; cartilage differentiation; wound healing;
XX neuron growth; bone fracture; osteoporosis; osteopenia; arthritis;
XX sarcopenia; periodontal disease; tissue atrophy; endocrine disorder;
XX muscle loss; immobility; bone density; mutant; muteln.

```

XX OS Homo sapiens.
XX OS Synthetic.
XX FH Key Location/Qualifiers
XX FT Misc-difference 276
XX FT /note= "Wild type Asn substituted with Asp"
XX PN W0200189450-A2.
XX PD 29-NOV-2001.
XX PE 08-MAY-2001; 2001WO-US11755.
XX PR 19-MAY-2000; 2000US-205424P.
XX PR 11-JAN-2001; 2001US-261071P.
XX PR 11-JAN-2001; 2001US-261076P.
XX PA (ELIL ) LILLY & CO ELI.
XX PI Beals JM, Gonzalez-Dewhilt PA, Hammond LJ, Lu J, Na S, Su EW;
XX PI Witcher DR, Wroblewski VJ;
XX DR WPI: 2002-083040/11.
XX PT Analog of a platelet-derived growth factor homolog, LP85 useful for
XX PT treating osteoporosis, arthritis, sarcopenia, wounds, has one or more
XX PT amino acid substitutions which destroy the tripeptidyl sequence of
XX PT native LP85
XX PS Claim 15b; Page -: 117pp; English.
XX CC The present invention relates to LP85, an analogue of platelet-derived
XX CC growth factor (PDGF) homologue. Sequences of the invention are useful
XX CC for the manufacture of a medicament for treating musculoskeletal disorder
XX CC (MSD) which include promoting bone growth, cartilage differentiation and
XX CC function, wound healing, neuron growth, preventing cartilage degradation
XX CC or neuronal degeneration. They are useful for treating bone fractures,
XX CC osteoporosis, sarcopenia, arthritis, sarcopenia, periodontal disease,
XX CC tissue atrophy, traumatised connective tissues, grafted connective
XX CC tissues and/or transplanted organs, bone or muscle loss due to
XX CC malignancy, endocrine disorders and immobility. They are also used
XX CC for prophylactically increasing or maintaining bone density in a
XX CC mammal. The present sequence is human LP85 mutant protein (N276D).
XX CC Note: This sequence is not shown in the specification but is derived
XX CC from human LP85 protein shown in pages 109-110 of the specification
XX CC (AAE15819).
XX SQ Sequence 370 AA;
XX
Query Match 99.7%; Score 1989; DB 23; Length 370;
Best Local Similarity 99.7%; Pred. No. 3,5e-188;
Matches 369; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

```

```

QY 301 RCGNCGCGTYNMRSCTGNSGTYKYYHEVLOFEFGHTKRGRKATMALNDIOJDHHERC 360
DB 301 RCGNCGCGTYNMRSCTGNSGTYKYYHEVLOFEFGHTKRGRKATMALNDIOJDHHERC 360
QY 361 DCICSSRPPR 370
DB 361 DCICSSRPPR 370
RESULT 15
AAE15849
ID AAE15849 standard; Protein: 370 AA.
AC AAE15849;
XX 26-MAR-2002 (first entry)
XX
XX Human LP85 mutant protein (N276S).
XX
XX LP85; platelet-derived growth factor; PDGF; antiinflammatory; vulnerrary;
XX KM osteopathic; neuroprotective; tranquilliser; musculoskeletal disorder;
XX KM MSD; therapy; bone growth; cartilage differentiation; wound healing;
XX KM neuron growth; bone fracture; osteoporosis; osteopenia; arthritis;
XX KM sarcopenia; periodontal disease; tissue atrophy; endocrine disorder;
XX KM muscle loss; immobility; bone density; mutant; muteln.
XX
XX Homo sapiens.
XX OS Synthetic.
XX FH Key Location/Qualifiers
XX FT Misc-difference 276
XX FT /note= "Wild type Asn substituted with Ser"
XX FT W0200189450-A2.
XX PD 29-NOV-2001.
XX PE 08-MAY-2001; 2001WO-US11755.
XX PR 19-MAY-2000; 2000US-205424P.
XX PR 11-JAN-2001; 2001US-261071P.
XX PR 11-JAN-2001; 2001US-261076P.
XX PA (ELIL ) LILLY & CO ELI.
XX PI Beals JM, Gonzalez-Dewhilt PA, Hammond LJ, Lu J, Na S, Su EW;
XX PI Witcher DR, Wroblewski VJ;
XX DR WPI: 2002-083040/11.
XX PT Analog of a platelet-derived growth factor homolog, LP85 useful for
XX PT treating osteoporosis, arthritis, sarcopenia, wounds, has one or more
XX PT amino acid substitutions which destroy the tripeptidyl sequence of
XX PT native LP85
XX PS Claim 15f; Page -: 117pp; English.
XX CC The present invention relates to LP85, an analogue of platelet-derived
XX CC growth factor (PDGF) homologue. Sequences of the invention are useful
XX CC for the manufacture of a medicament for treating musculoskeletal disorder
XX CC (MSD) which include promoting bone growth, cartilage differentiation and
XX CC function, wound healing, neuron growth, preventing cartilage degradation
XX CC or neuronal degeneration. They are useful for treating bone fractures,
XX CC osteoporosis, sarcopenia, arthritis, sarcopenia, periodontal disease,
XX CC tissue atrophy, traumatised connective tissues, grafted connective
XX CC tissues and/or transplanted organs, bone or muscle loss due to
XX CC malignancy, endocrine disorders and immobility. They are also used
XX CC for prophylactically increasing or maintaining bone density in a
XX CC mammal. The present sequence is human LP85 mutant protein (N276S).
XX CC Note: This sequence is not shown in the specification but is derived
XX CC from human LP85 protein shown in pages 109-110 of the specification
XX CC (AAE15819).

```

xx Sequence 370 Aa;

Query Match 99.7%; Score 1989; DB 23; Length 370;
 Best Local Similarity 99.7%; Pred. No. 3.5e-188;
 Matches 369; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy	1	MHRLIFVYTLICANFCSCDTSATPOSASIKALRNANLRDESNHLDLYRDEFTQVKG	60
Db	1	MHRLIFVYTLICANFCSCDTSATPOSASIKALRNANLRDESNHLDLYRDEFTQVKG	60
Qy	61	NGYVSPRPNSYPNNLLTWRLHSQENTRIQLVFNQGLEAEENDICRYDFVEVEDIS	120
Db	61	NGYVSPRPNSYPNNLLTWRLHSQENTRIQLVFNQGLEAEENDICRYDFVEVEDIS	120
Qy	121	ETSTIIRGRCGCHKKEVPPRIKSRTOIKITFKSDYFVAKPGFKIYSLLEDFOPAASE	180
Db	121	ETSTIIRGRCGCHKKEVPPRIKSRTOIKITFKSDYFVAKPGFKIYSLLEDFOPAASE	180
Qy	181	TNMEVTSISISGVSNPSVTDPPTLIADALDKKIAEPTVEDLKYFNPESMQEDLENNY	240
Db	181	TNMEVTSISISGVSNPSVTDPPTLIADALDKKIAEPTVEDLKYFNPESMQEDLENNY	240
Qy	241	LDPTRYGRSYHDKRSKYDLRLNDARKYSCTPRNYSVINIREELKIANVVFPRCLVQ	300
Db	241	LDPTRYGRSYHDKRSKYDLRLNDARKYSCTPRNYSVINIREELKIANVVFPRCLVQ	300
Qy	301	RCGNCGGGTVMNSCTCNSGKYKYEVLQFEFGHIKRRGAKTMALVDIQLDHHERC	360
Db	301	RCGNCGGGTVMNSCTCNSGKYKYEVLQFEFGHIKRRGAKTMALVDIQLDHHERC	360
Qy	361	DCICSSRPPR 370	
Db	361	DCICSSRPPR 370	

Search completed: June 12, 2003, 15:29:58
 Job time : 51 secs

GenCore version 5.1.6
Copyright (c) 1993 - 2003 CompuGen Ltd.

OM protein - protein search, using sw model

Run on: June 12, 2003, 15:28:35 ; Search time 26 Seconds
(without alignments)
418.711 Million cell updates/sec

Title: US-09-662-783-2
Perfect score: 1994
Sequence: 1 MHRLLFYVTLICANFCSCRD.....DIDLHHERCDICSSRPPR 370

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 262574 seqs, 29422922 residues

Total number of hits satisfying chosen parameters: 262574

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%

Listing first 45 summaries

Database : Issued_Patents_AA:*
1: /cgn2_6/p/ptodata/1/1aa/5A_COMB.pep:*
2: /cgn2_6/p/ptodata/1/1aa/5B_COMB.pep:*
3: /cgn2_6/p/ptodata/1/1aa/6A_COMB.pep:*
4: /cgn2_6/p/ptodata/1/1aa/6B_COMB.pep:*
5: /cgn2_6/p/ptodata/1/1aa/PTUS_COMB.pep:*
6: /cgn2_6/p/ptodata/1/1aa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	1994	100.0	370	US-09-457-066-37	Sequence 37, Appl
2	1994	100.0	370	US-09-540-224-2	Sequence 2, Appl
3	1737	87.1	370	US-09-540-224-4	Sequence 4, Appl
4	752	37.7	345	US-09-457-066-43	Sequence 43, Appl
5	742.5	37.2	345	US-09-040-220D-2	Sequence 2, Appl
6	742.5	37.2	345	US-09-457-066-2	Sequence 2, Appl
7	742.5	37.2	345	US-09-265-686-2	Sequence 2, Appl
8	742.5	37.2	345	US-09-540-224-5	Sequence 5, Appl
9	187.5	9.4	788	US-08-572-225-1	Sequence 1, Appl
10	186.5	9.4	730	US-08-872-757-2	Sequence 2, Appl
11	180.5	9.1	101	US-09-374-135-6	Sequence 6, Appl
12	174.5	8.8	986	US-08-872-757-4	Sequence 4, Appl
13	172.5	8.7	591	US-08-991-408-4	Sequence 4, Appl
14	172.5	8.7	591	US-09-432-473-4	Sequence 4, Appl
15	172.5	8.7	1013	US-08-866-650-5	Sequence 5, Appl
16	172.5	8.7	1013	US-09-021-287-5	Sequence 5, Appl
17	172.5	8.7	1013	US-08-991-408-2	Sequence 2, Appl
18	172.5	8.7	1013	US-09-240-473-5	Sequence 5, Appl
19	172.5	8.7	1013	US-09-432-473-2	Sequence 2, Appl
20	167.5	8.4	103	US-09-374-135-5	Sequence 5, Appl
21	167	8.4	922	US-09-116-473-4	Sequence 4, Appl
22	166.5	8.4	1013	US-08-866-650-3	Sequence 3, Appl
23	166.5	8.4	1013	US-09-021-287-3	Sequence 3, Appl
24	166.5	8.4	1013	US-09-240-473-3	Sequence 3, Appl
25	164	8.2	923	US-08-936-135-6	Sequence 6, Appl
26	159	8.0	909	US-08-936-135-18	Sequence 18, Appl
27	159	8.0	926	US-08-936-135-20	Sequence 20, Appl

28	157	7.9	901	US-08-936-135-22	Sequence 22, Appl
29	157	7.9	906	US-08-936-135-24	Sequence 24, Appl
30	157	7.9	909	US-08-936-135-8	Sequence 8, Appl
31	157	7.9	909	US-08-936-135-10	Sequence 10, Appl
32	157	7.9	914	US-08-936-135-12	Sequence 12, Appl
33	157	7.9	925	US-09-116-473-2	Sequence 2, Appl
34	157	7.9	926	US-08-936-135-14	Sequence 14, Appl
35	157	7.9	931	US-08-936-135-16	Sequence 16, Appl
36	144	7.2	1785	US-09-341-587-3	Sequence 3, Appl
37	142	7.1	101	US-09-374-135-4	Sequence 4, Appl
38	138	6.9	25	US-09-540-224-8	Sequence 8, Appl
39	136.5	6.8	449	US-08-839-008-9	Sequence 9, Appl
40	135.5	6.8	102	US-09-374-135-7	Sequence 7, Appl
41	131	6.6	666	US-09-341-587-1	Sequence 1, Appl
42	130.5	6.5	401	US-08-839-008-5	Sequence 5, Appl
43	130.5	6.5	415	US-09-032-523-2	Sequence 2, Appl
44	130.5	6.5	449	US-08-839-008-2	Sequence 2, Appl
45	130.5	6.5	468	US-08-839-008-7	Sequence 7, Appl

ALIGNMENTS

RESULT 1
US-09-457-066-37
Sequence 37, Application US/09457066
Patent No. 6432673
GENERAL INFORMATION:
APPLICANT: Gao, Zhen
APPLICANT: Hart, Charles E.
APPLICANT: Piddington, Christopher S.
APPLICANT: Sheppard, Paul O.
APPLICANT: Shomaker, Kimberly E.
APPLICANT: Gilbertson, Debra G.
APPLICANT: West, James W.
FILE REFERENCE: 98-60
CURRENT APPLICATION NUMBER: US/09/457,066
CURRENT FILING DATE: 1999-12-07
NUMBER OF SEQ ID NOS: 50
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 37
LENGTH: 370
TYPE: PRT
ORGANISM: Homo sapiens
US-09-457-066-37

Query Match
Best local similarity 100.0%; Score 1994; DB 4; Length 370;
Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MHRLLFYVTLICANFCSCROTSATPOSASIKALNANLRDSEHMLDLYRDETQYKG 60
Db 1 MHRLLFYVTLICANFCSCROTSATPOSASIKALNANLRDSEHMLDLYRDETQYKG 60
QY 61 NGYQSPREFPSTYRNLLTWRLHSENTRIQVFNQFGLAEANDICRYFEVEDIS 120
Db 61 NGYQSPREFPSTYRNLLTWRLHSENTRIQVFNQFGLAEANDICRYFEVEDIS 120
QY 121 ETSITIRGRCWGKHEVPRIKSRTOIKITFKSDYEVAKFGFIYYSLEDFOPAAASE 180
Db 121 ETSITIRGRCWGKHEVPRIKSRTOIKITFKSDYEVAKFGFIYYSLEDFOPAAASE 180
QY 181 TNMESVYSSISGVNSYVDPLLDALDKKTAEDPTVBDLKYFNPESWQDLEMY 240
Db 181 TNMESVYSSISGVNSYVDPLLDALDKKTAEDPTVBDLKYFNPESWQDLEMY 240
QY 241 LDTRYGRGRSHDKRSKYVDLRLNDADKRYSCPTPNTSVNREELKLANVFFPRCLLYQ 300
Db 241 LDTRYGRGRSHDKRSKYVDLRLNDADKRYSCPTPNTSVNREELKLANVFFPRCLLYQ 300
QY 301 RCGNCGGCTVNMWSTCTNSGKTVKYYHEVLQEPFGHKKRGRKATMALVDIQLDHHERC 360
Db 301 RCGNCGGCTVNMWSTCTNSGKTVKYYHEVLQEPFGHKKRGRKATMALVDIQLDHHERC 360

Db 301 RCGNGCGGVNMRSCSTGSKTYKTYHEVLQEPFGHKKRGAKTALVDIQLDHHERC 360
 QY 361 DCICSSRPR 370
 Db 361 DCICSSRPR 370

RESULT 2

US-09-540-224-2
 ; Sequence 2, Application US/09540224
 ; Patent No. 6468543
 ; GENERAL INFORMATION:
 ; APPLICANT: Gilbertson, Debra G.
 ; TITLE OF INVENTION: METHODS FOR PROMOTING GROWTH OF BONE,
 ; FILE REFERENCE: 00-28
 ; CURRENT APPLICATION NUMBER: US/09/540,224
 ; EARLIER FILING DATE: 2000-03-31
 ; EARLIER APPLICATION NUMBER: US 60/180,169
 ; NUMBER OF SEQ ID NOS: 9
 ; SOFTWARE: FastSeq for Windows Version 3.0
 ; SEQ ID NO 2
 ; LENGTH: 370
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 US-09-540-224-2

Query Match 100.0%; Score 1994; DB 4; Length 370;
 Best Local Similarity 100.0%; Pred. No. 4, 7e-198;
 Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MHRLLIVYLLICANFCSCRDTSATPOSASIKALRNANLRDESNHLLDLYRDETIQYKG 60
 Db 1 MHRLLIVYLLICANFCSCRDTSATPOSASIKALRNANLRDESNHLLDLYRDETIQYKG 60
 QY 61 NGVQSPRPNSYPNNLLTWRLHSGENTRIQLVFNQFGLAEANDICRYDFVEVEDIS 120
 Db 61 NGVQSPRPNSYPNNLLTWRLHSGENTRIQLVFNQFGLAEANDICRYDFVEVEDIS 120
 QY 121 ESTTIIRGRCGHEKVEPPRIKSRNTQIKITFKSDDFVAKPGFKIYSLLEDFQPAASE 180
 Db 121 ESTTIIRGRCGHEKVEPPRIKSRNTQIKITFKSDDFVAKPGFKIYSLLEDFQPAASE 180
 QY 181 TWNESTVSSISGYSYSPSTVDTPTLLADALDKKIAEFDTVEDLLKYNPESMOEDLENMY 240
 Db 181 TWNESTVSSISGYSYSPSTVDTPTLLADALDKKIAEFDTVEDLLKYNPESMOEDLENMY 240
 QY 241 LDTPRYGRSYHDKRSKVDLRLNDADAKRYSCPTPRNYSVIRELKLNAVFFPRCLLYQ 300
 Db 241 LDTPRYGRSYHDKRSKVDLRLNDADAKRYSCPTPRNYSVIRELKLNAVFFPRCLLYQ 300
 QY 301 RCGNGCGGVNMRSCSTGSKTYKTYHEVLQEPFGHKKRGAKTALVDIQLDHHERC 360
 Db 301 RCGNGCGGVNMRSCSTGSKTYKTYHEVLQEPFGHKKRGAKTALVDIQLDHHERC 360
 QY 361 DCICSSRPR 370
 Db 361 DCICSSRPR 370

RESULT 3

US-09-540-224-4
 ; Sequence 4, Application US/09540224
 ; Patent No. 6468543
 ; GENERAL INFORMATION:
 ; APPLICANT: Gilbertson, Debra G.
 ; TITLE OF INVENTION: METHODS FOR PROMOTING GROWTH OF BONE,
 ; FILE REFERENCE: 00-28
 ; CURRENT APPLICATION NUMBER: US/09/540,224

; CURRENT FILING DATE: 2000-03-31
 ; EARLIER APPLICATION NUMBER: US 60/180,169
 ; EARLIER FILING DATE: 2000-02-04
 ; NUMBER OF SEQ ID NOS: 9
 ; SOFTWARE: FastSeq for Windows Version 3.0
 ; SEQ ID NO 4
 ; LENGTH: 370
 ; TYPE: PRT
 ; ORGANISM: Mus musculus
 US-09-540-224-4

Query Match 87.1%; Score 1737; DB 4; Length 370;
 Best Local Similarity 85.1%; Pred. No. 2e-171;
 Matches 315; Conservative 25; Mismatches 30; Indels 0; Gaps 0;

QY 1 MHRLLIVYLLICANFCSCRDTSATPOSASIKALRNANLRDESNHLLDLYRDETIQYKG 60
 Db 1 MHRLLIVYLLICANFCSCRDTSATPOSASIKALRNANLRDESNHLLDLYRDETIQYKG 60
 QY 61 NGVQSPRPNSYPNNLLTWRLHSGENTRIQLVFNQFGLAEANDICRYDFVEVEDIS 120
 Db 61 NGVQSPRPNSYPNNLLTWRLHSGENTRIQLVFNQFGLAEANDICRYDFVEVEDIS 120
 QY 121 ESTTIIRGRCGHEKVEPPRIKSRNTQIKITFKSDDFVAKPGFKIYSLLEDFQPAASE 180
 Db 121 ESTTIIRGRCGHEKVEPPRIKSRNTQIKITFKSDDFVAKPGFKIYSLLEDFQPAASE 180
 QY 181 TWNESTVSSISGYSYSPSTVDTPTLLADALDKKIAEFDTVEDLLKYNPESMOEDLENMY 240
 Db 181 TWNESTVSSISGYSYSPSTVDTPTLLADALDKKIAEFDTVEDLLKYNPESMOEDLENMY 240
 QY 241 LDTPRYGRSYHDKRSKVDLRLNDADAKRYSCPTPRNYSVIRELKLNAVFFPRCLLYQ 300
 Db 241 LDTPRYGRSYHDKRSKVDLRLNDADAKRYSCPTPRNYSVIRELKLNAVFFPRCLLYQ 300
 QY 301 RCGNGCGGVNMRSCSTGSKTYKTYHEVLQEPFGHKKRGAKTALVDIQLDHHERC 360
 Db 301 RCGNGCGGVNMRSCSTGSKTYKTYHEVLQEPFGHKKRGAKTALVDIQLDHHERC 360
 QY 361 DCICSSRPR 370
 Db 361 DCICSSRPR 370

RESULT 4

US-09-457-066-43
 ; Sequence 43, Application US/09457066
 ; Patent No. 6432673
 ; GENERAL INFORMATION:
 ; APPLICANT: Gao, Zeren
 ; APPLICANT: Hart, Charles E.
 ; APPLICANT: Piddington, Christopher S.
 ; APPLICANT: Sheppard, Paul O.
 ; APPLICANT: Shoemaker, Kimberly E.
 ; APPLICANT: Gilbertson, Debra G.
 ; TITLE OF INVENTION: GROWTH FACTOR HOMOLOG ZVEGF3
 ; FILE REFERENCE: 98-60
 ; CURRENT APPLICATION NUMBER: US/09/457,066
 ; CURRENT FILING DATE: 1999-12-07
 ; NUMBER OF SEQ ID NOS: 50
 ; SOFTWARE: FastSeq for Windows Version 3.0
 ; SEQ ID NO 43
 ; LENGTH: 345
 ; TYPE: PRT
 ; ORGANISM: Mus musculus
 US-09-457-066-43

Query Match 37.7%; Score 752; DB 4; Length 345;
 Best Local Similarity 45.3%; Pred. No. 1.8e-69;
 Matches 148; Conservative 59; Mismatches 92; Indels 28; Gaps 9;

QY 42 ESNHLLDLYRDETIQYKNGYVQSPRPNSYPNNLLTWRLHSGENTRIQLVFNQFGLAEANDICRYDFVEVEDIS 100

```
Db      37 EQNGVOD-PHREVRVITSGNSIHSFKFPHYPRNVLVRLVAVDENVRQLQLEFDEBRFG 95
Qy      101 LEEAENDICRYDPEVEVDISETSTIIRGRCGKEVPRPKISRNQIKITFKSDIYVAK 160
Db      96 LEEPEDICRYDPEVEVEPDSGVL--GRMGSGTGVGKQISKGNHIRIRFVDEYEPSE 153
Qy      161 PGFKIYSLLEDFOPAASETNMESVTSISGVSPSYTDP--TLADLADKKIAEDT 219
Db      154 PGECIHSTI--MPQYETI-----SPVLPSPSLDLNNAVTAFTST 195
Qy      220 VEDLLKYNPESQEDLENNYLDTPRYGRSY-HDRKSK-VDLDRLNDKARYSCTPRNY 277
Db      196 LEEILRYLEPDRMQVDLSLYKPTWQLGKAFYLGKSKSVNLMLKEEYVYSCPTPNE 255
Qy      278 SVNIRELKLANYVEFRCLLYVORCGNGCGGVNMSCTCNSKTYKTHYEVQFEPBGH 337
Db      256 SVSIRELKRDTITFWPCCLLYKRCGNCACCLHNCOCQVPRKTYKHYEVLQLRP-- 313
Qy      338 IKRGRAKTMALVDIOLDHHERCDIC 364
Db      314 -KIGVKGILKSLTDVVALEHHECDVC 339
```

```
RESULT 5
US-09-040-220D-2
; Sequence 2, Application US/09040220D
; Patent No. 6391311
; GENERAL INFORMATION:
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Kuo, Sophia S.
; TITLE OF INVENTION: NOVEL POLYPEPTIDES HAVING HOMOLOGY TO VASCULAR
; TITLE OF INVENTION: ENDOCELLULAR CELL GROWTH FACTOR AND BONE MORPHOGENETIC
; TITLE OF INVENTION: PROTEIN 1 AND NUCLEIC ACIDS ENCODING SAME, THEIR USES,
; FILE REFERENCE: P1122
; CURRENT APPLICATION NUMBER: US/09/040.220D
; CURRENT FILING DATE: 1998-03-17
; NUMBER OF SEQ ID NOS: 8
; SEQ ID NO 2
; LENGTH: 345
; TYPE: PRT
; ORGANISM: Human
US-09-040-220D-2
```

```
Query Match      37.2%; Score 742.5; DB 4; Length 345;
Best Local Similarity 43.6%; Pred. No. 1.8e-68;
Matches 159; Conservative 59; Mismatches 114; Indels 33; Gaps 10;
```

```
Qy      5 IFVYTLICANFCSCROTSATPQASISIKALRNANLRDESNHLTDLYRDETIOYKNGYV 64
Db      3 LFGILLITSLAGOROGTOAESNLSSKFQSSN--KEQNGVOD-PQHERITIVSTNGSI 58
Qy      65 QSPRPNSYPNLLITWRKLS-QENTRIQVFDNOFGLEAENDICRYDVEVEDISETS 123
Db      59 HSPRPPTYPRNTLVWRLVAVEENWVQLTFDERFGLDEDEDICRYDVEVEPDSGT 118
Qy      124 TIIRGRCGKEVPRPKISRNQIKITFKSDIYVAKPGFKIYSL--LEDFOPAASETN 182
Db      119 --ILGRMGSGTGVGKQISKGNQIRIRFVSEYEPSEPGFCIHYNIVAPQTEAV---- 171
Qy      183 WESTSSISGVSPSYTDP--LIADLADKKIAEDTVDLLKYFNPESQEDLENNYL 241
Db      172 -----SPVLPSPSALPLDLNNAITAFSTLEDLIRYLEERWQDLEDLYR 217
Qy      242 DTPRYGRSY-HDRKSK-VDLDRLNDKARYSCTPRNYSVIRELKLANYVEFRCLLY 299
Db      218 PTWQLGKAFYFGKRSRVVDLNLTEEVRLYSCTPRNSVSIRELKRKTDTITFWPCCLLY 277
Qy      300 QRCGNGCGGTIVNRSCTCNSGKTYKTHYEVQFEPGHIKRRGAKTMALVDIOLDHHER 359
Db      278 KRCGNCACCLHNCNEOCQVSKTYKHYEVLQLRP--KIGVKGILKSLTDVVALEHHEE 334
```

```
Qy      360 CDCIC 364
Db      335 CDCVC 339
```

```
RESULT 6
US-09-457-066-2
; Sequence 2, Application US/09457066
; Patent No. 6432673
; GENERAL INFORMATION:
; APPLICANT: Gao, Zeren
; APPLICANT: Hart, Charles E.
; APPLICANT: Piddington, Christopher S.
; APPLICANT: Sheppard, Paul O.
; APPLICANT: Shoemaker, Kimberly E.
; APPLICANT: Gilbertson, Debra G.
; TITLE OF INVENTION: GROWTH FACTOR HOMOLOGY ZVEGF3
; FILE REFERENCE: 98-60
; CURRENT APPLICATION NUMBER: US/09/457.066
; CURRENT FILING DATE: 1999-12-07
; NUMBER OF SEQ ID NOS: 50
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 2
; LENGTH: 345
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-457-066-2
```

```
Query Match      37.2%; Score 742.5; DB 4; Length 345;
Best Local Similarity 43.6%; Pred. No. 1.8e-68;
Matches 159; Conservative 59; Mismatches 114; Indels 33; Gaps 10;
```

```
Qy      5 IFVYTLICANFCSCROTSATPQASISIKALRNANLRDESNHLTDLYRDETIOYKNGYV 64
Db      3 LFGILLITSLAGOROGTOAESNLSSKFQSSN--KEQNGVOD-PQHERITIVSTNGSI 58
Qy      65 QSPRPNSYPNLLITWRKLS-QENTRIQVFDNOFGLEAENDICRYDVEVEDISETS 123
Db      59 HSPRPPTYPRNTLVWRLVAVEENWVQLTFDERFGLDEDEDICRYDVEVEPDSGT 118
Qy      124 TIIRGRCGKEVPRPKISRNQIKITFKSDIYVAKPGFKIYSL--LEDFOPAASETN 182
Db      119 --ILGRMGSGTGVGKQISKGNQIRIRFVSEYEPSEPGFCIHYNIVAPQTEAV---- 171
Qy      183 WESTSSISGVSPSYTDP--LIADLADKKIAEDTVDLLKYFNPESQEDLENNYL 241
Db      172 -----SPVLPSPSALPLDLNNAITAFSTLEDLIRYLEERWQDLEDLYR 217
Qy      242 DTPRYGRSY-HDRKSK-VDLDRLNDKARYSCTPRNYSVIRELKLANYVEFRCLLY 299
Db      218 PTWQLGKAFYFGKRSRVVDLNLTEEVRLYSCTPRNSVSIRELKRKTDTITFWPCCLLY 277
Qy      300 QRCGNGCGGTIVNRSCTCNSGKTYKTHYEVQFEPGHIKRRGAKTMALVDIOLDHHER 359
Db      278 KRCGNCACCLHNCNEOCQVSKTYKHYEVLQLRP--KIGVKGILKSLTDVVALEHHEE 334
Qy      360 CDCIC 364
Db      335 CDCVC 339
```

```
RESULT 7
US-09-265-686-2
; Sequence 2, Application US/09265686
; Patent No. 6455283
; GENERAL INFORMATION:
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Kuo, Sophia S.
; TITLE OF INVENTION: POLYPEPTIDES HOMOLOGOUS TO VEGF AND BMP1
; FILE REFERENCE: P1122P2
; CURRENT APPLICATION NUMBER: US/09/265.686
; CURRENT FILING DATE: 1999-03-10
```

PRIOR APPLICATION NUMBER: US 09/040,220
PRIOR FILING DATE: 1998-03-17
PRIOR APPLICATION NUMBER: US 09/184,216
PRIOR FILING DATE: 1998-11-02
NUMBER OF SEQ ID NOS: 8
SEQ ID NO: 2
LENGTH: 345
TYPE: PRT
ORGANISM: Human
US-09-265-686-2

Query Match 37.2%; Score 742.5; DB 4; Length 345;
Best Local Similarity 43.6%; Pred. No. 1.8e-68;
Matches 159; Conservative 59; Mismatches 114; Indels 33; Gaps 10;

QY 5 IFVYTLICANFCSCRTSATPOSASIKALNANLRDESNHLDLVRDETIOYKNGYV 64
DB 3 LFEULLITSLAQROGTOAESNLSSKFOFSSN---KEONGVOD-POHERITIVSTNGSI 58
QY 65 QSPRPNSYPNNLLTWRLHS-OENRIQLVFNQGLEAENDICRYDEVEEDISETS 123
DB 59 HSPRFHTYPRNTVLVWRLVAEENWVQLTFDERFGLDEPDICRYDEVEEEDSDGT 118
QY 124 TIIRGMCCKEYPPRIKSTNOIKTFKSDDYFVAKPGFKIYSL-LEDFOPAAASETN 182
DB 119 --ILGRWCGSGTVPGRKQISGNQIRIFVSDYFSPSGFCHYNIVMPQTEAV----- 171
QY 183 WESVTSISGSVNSPSTPDT-LIADALDKIAEFDVLDLKYPNESMODLEMYL 241
DB 172 -----SPSVLPSPSALPLDLNNAITASTLEDLIRYLEPERMQLDELYR 217
QY 242 DTPRYGRSY-HDRKSK-VDLRLNDAKRYCTPRNYSVIREELKLANVFEPRCLIV 299
DB 218 PTMQLGKAFAVFGKRSVVDNLNLTBEVRLSCTPRNYSVIREELKRTDTIWPGLIV 277
QY 300 QRCGNCGGCTVWNRSCNCGTKYKHEVLOFEPGHIRRRRAKTMALVDIQLDHER 359
DB 278 KRCGNCACCLHNCNEOCVPSKYTKYHEVLOLRP---KTGYRGLHKSILVDALHEHEE 334
QY 360 CDCVC 364
DB 335 CDCVC 339

RESULT 8
US-09-540-224-5
Sequence 5, Application US/09540224
Patent No. 6468543
GENERAL INFORMATION:
APPLICANT: Gilbertson, Debra G.
APPLICANT: Hart, Charles E.
TITLE OF INVENTION: METHODS FOR PROMOTING GROWTH OF BONE,
TITLE OF INVENTION: LIGAMENT AND CARTILAGE USING ZVEGH4
FILE REFERENCE: 00-28
CURRENT APPLICATION NUMBER: US/09/540,224
CURRENT FILING DATE: 2000-03-31
EARLIER APPLICATION NUMBER: US 60/180,169
EARLIER FILING DATE: 2000-02-04
NUMBER OF SEQ ID NOS: 9
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO: 5
LENGTH: 345
TYPE: PRT
ORGANISM: Homo sapiens
US-09-540-224-5

Query Match 37.2%; Score 742.5; DB 4; Length 345;
Best Local Similarity 43.6%; Pred. No. 1.8e-68;
Matches 159; Conservative 59; Mismatches 114; Indels 33; Gaps 10;

QY 5 IFVYTLICANFCSCRTSATPOSASIKALNANLRDESNHLDLVRDETIOYKNGYV 64
DB 3 LFEULLITSLAQROGTOAESNLSSKFOFSSN---KEONGVOD-POHERITIVSTNGSI 58

QY 65 QSPRPNSYPNNLLTWRLHS-OENRIQLVFNQGLEAENDICRYDEVEEDISETS 123
DB 59 HSPRFHTYPRNTVLVWRLVAEENWVQLTFDERFGLDEPDICRYDEVEEEDSDGT 118
QY 124 TIIRGMCCKEYPPRIKSTNOIKTFKSDDYFVAKPGFKIYSL-LEDFOPAAASETN 182
DB 119 --ILGRWCGSGTVPGRKQISGNQIRIFVSDYFSPSGFCHYNIVMPQTEAV----- 171
QY 183 WESVTSISGSVNSPSTPDT-LIADALDKIAEFDVLDLKYPNESMODLEMYL 241
DB 172 -----SPSVLPSPSALPLDLNNAITASTLEDLIRYLEPERMQLDELYR 217
QY 242 DTPRYGRSY-HDRKSK-VDLRLNDAKRYCTPRNYSVIREELKLANVFEPRCLIV 299
DB 218 PTMQLGKAFAVFGKRSVVDNLNLTBEVRLSCTPRNYSVIREELKRTDTIWPGLIV 277
QY 300 QRCGNCGGCTVWNRSCNCGTKYKHEVLOFEPGHIRRRRAKTMALVDIQLDHER 359
DB 278 KRCGNCACCLHNCNEOCVPSKYTKYHEVLOLRP---KTGYRGLHKSILVDALHEHEE 334
QY 360 CDCVC 364
DB 335 CDCVC 339

RESULT 9
US-08-572-225-1
Sequence 1, Application US/08572225
Patent No. 5807981
GENERAL INFORMATION:
APPLICANT: Prockop, Darwin J.
APPLICANT: Hojima, Yoshio
APPLICANT: Li, Shi-Wu
APPLICANT: Stierou, Alexander
APPLICANT: Brenner, Mitch
TITLE OF INVENTION: RECOMBINANT C-PROTEINASE AND ITS USE FOR
TITLE OF INVENTION: DRUG DEVELOPMENT FOR THE TREATMENT OF DISEASE
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: Penile & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/572,225
FILING DATE: 13-DEC-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Halluin, Albert P.
REGISTRATION NUMBER: 25,227
REFERENCE/DOCKET NUMBER: 8389-031
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-3660
TELEFAX: 415-854-3694
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 788 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: protein
US-08-572-225-1

Query Match 9.4%; Score 187.5; DB 1; Length 788;

Best Local Similarity 39.4%; Pred. No. 2e-10;
Matches 43; Conservative 18; Mismatches 43; Indels 5; Gaps 3;

OY 59 KGNQVSPRPNSYPRNLLTWRHLSQENTRIQLVFDNPGLEAEANDICRYFVEVED 118
DB 399 KLNQSTSPGPKPEYPPKNCIMQVAPTOYRISLOPD---FFETEGNDVCKYDFVEVRS 455
OY 119 ISESTIRGRCGKHEVPPRIKSRTOIKITFKSDDYFAKPGFKIYY 167
DB 456 GLTADSKLHGKFCG-SEKPEVITTSQYNNMRYEFSKSDN-TVSKGKFAHF 502

RESULT 10

US-08-872-757-2
Sequence 2, Application US/08872757
Patent No. 6258584
GENERAL INFORMATION:
APPLICANT: Prockop, Darwin J.
APPLICANT: Hojima, Yoshio
APPLICANT: Li, Shi-Wu
APPLICANT: Steron, Aleksander
TITLE OF INVENTION: RECOMBINANT C-PROTEINASE AND
TITLE OF INVENTION: PROCESSES; METHODS AND USES THEREOF
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESSES:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/872,757
FILING DATE: 10-JUN-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/609,187
FILING DATE: 01-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Halluin, Albert P.
REGISTRATION NUMBER: 25,227
REFERENCE/DOCKET NUMBER: 8389-028-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-3660
TELEFAX: 415-854-3694
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO. 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 730 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-872-757-2

Query Match 9.4%; Score 186.5; DB 4; Length 730;
Best Local Similarity 39.4%; Pred. No. 2.3e-10;
Matches 43; Conservative 18; Mismatches 43; Indels 5; Gaps 3;

OY 59 KGNQVSPRPNSYPRNLLTWRHLSQENTRIQLVFDNPGLEAEANDICRYFVEVED 118
DB 597 KLNQSTSPGPKPEYPPKNCIMQVAPTOYRISLOPD---FFETEGNDVCKYDFVEVRS 653
OY 119 ISESTIRGRCGKHEVPPRIKSRTOIKITFKSDDYFAKPGFKIYY 167
DB 654 GLTADSKLHGKFCG-SEKPEVITTSQYNNMRYEFSKSDN-TVSKGKFAHF 700

RESULT 11

US-09-374-135-6
Sequence 6, Application US/09374135
Patent No. 6277972

GENERAL INFORMATION:
APPLICANT: Afari, Daniel E.
APPLICANT: Hubert, Rene S.
APPLICANT: Leong, Kahan
APPLICANT: Raitano, Arthur B.
APPLICANT: Saffran, Douglas C.
APPLICANT: Jakobovits, Aya
TITLE OF INVENTION: BPC-1: A SECRETED BRAIN-SPECIFIC PROTEIN EXPRESSED AND
SECRETED BY PROSTATE AND BLADDER CANCER CELLS
FILE REFERENCE: 1703-017 US1
CURRENT APPLICATION NUMBER: US/09/374,135
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: 60/095,982
PRIOR FILING DATE: 1998-08-10
NUMBER OF SEQ ID NOS: 20
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 6
LENGTH: 101
TYPE: PRT
ORGANISM: Mouse
US-09-374-135-6

Query Match 9.1%; Score 180.5; DB 4; Length 101;
Best Local Similarity 38.7%; Pred. No. 4e-11;
Matches 41; Conservative 18; Mismatches 42; Indels 5; Gaps 3;

OY 62 GYVSPRPNSYPRNLLTWRHLSQENTRIQLVFDNPGLEAEANDICRYFVEVEDISE 121
DB 1 GSITSPGPKPEYPPKNCIMQVAPTOYRISLOPD---FFETEGNDVCKYDFVEVRSGLT 57
OY 122 TSTIRGRCGKHEVPPRIKSRTOIKITFKSDDYFAKPGFKIYY 167
DB 58 ADSKLHGKFCG-SEKPEVITTSQYNNMRYEFSKSDN-TVSKGKFAHF 101

RESULT 12

US-08-872-757-4
Sequence 4, Application US/08872757
Patent No. 6258584
GENERAL INFORMATION:
APPLICANT: Prockop, Darwin J.
APPLICANT: Hojima, Yoshio
APPLICANT: Li, Shi-Wu
APPLICANT: Steron, Aleksander
TITLE OF INVENTION: RECOMBINANT C-PROTEINASE AND
TITLE OF INVENTION: PROCESSES; METHODS AND USES THEREOF
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESSES:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/872,757
FILING DATE: 10-JUN-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/609,187
FILING DATE: 01-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Halluin, Albert P.
REGISTRATION NUMBER: 25,227
REFERENCE/DOCKET NUMBER: 8389-028-999

TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-3660
TELEFAX: 415-854-3694
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 986 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-872-757-4

Query Match 8.8%; Score 174.5; DB 4; Length 986;
Best Local Similarity 37.6%; Pred. No. 6.5e-09;
Matches 41; Conservative 18; Mismatches 45; Indels 5; Gaps 3;

QY 59 KGNQVQSPRPSPYPRNLLITRLHSQENTRIQLVDFDNOFGLEAENDICRDEYVED 118
Db 597 KLNQGISRGMPREYEPNKNKCIQVAPTOYRISLQD---FFETEGNDYCKIDFEVRS 653
QY 119 ISETSTIRGRCMGHKEVPPRIKSRNQIKITFKSDDYFAKPGFKIYY 167
Db 654 GLTADSKLHGKFCG-SEKPEVITSQYNNMVEFEKSDN-TVSKKGKFAHF 700

RESULT 13
US-08-991-408-4
Sequence 4, Application US/08991408
Patent No. 6008017
GENERAL INFORMATION:
APPLICANT: ARLETH, ANTHONY J.
APPLICANT: WILLETTTE, ROBERT N.
APPLICANT: ELSHOURBAGY, NABIL A.
APPLICANT: LI, XIAOTONG
TITLE OF INVENTION: HUMAN CARDIAC/BRAIN TOLLDOID-LIKE
TITLE OF INVENTION: PROTEIN
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: RATNER & PRESTIA
STREET: P.O. BOX 980
CITY: VALLEY FORGE
STATE: PA
COUNTRY: USA
ZIP: 19482
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSRO for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/991,408
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/034,471
FILING DATE: 02-JAN-1997
ATTORNEY/AGENT INFORMATION:
NAME: PRESTIA, PAUL F
REGISTRATION NUMBER: 23,031
REFERENCE/DOCKET NUMBER: ATG-50038
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-407-0700
TELEFAX: 610-407-0701
TELEX: 846169
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 591 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-991-408-4

Query Match 8.7%; Score 172.5; DB 3; Length 591;
Best Local Similarity 36.7%; Pred. No. 4.6e-09;
Matches 40; Conservative 25; Mismatches 39; Indels 5; Gaps 4;

QY 59 KGNQVQSPRPSPYPRNLLITRLHSQENTRIQLVDFDNOFGLEAENDICRDEYVED 118
Db 202 KLNQGITTPGMPREYEPNKNKCIQVAPTOYRISVKE--FEELE--GNEVCKYDYEIWS 258
QY 119 ISETSTIRGRCMGHKEVPPRIKSRNQIKITFKSDDYFAKPGFKIYY 167
Db 259 GLSESKLHGKFCG-AEVEVITSQYNNMVEFEKSDN-TVSKKGKFAHF 305

RESULT 14
US-09-432-473-4
Sequence 4, Application US/09432473
Patent No. 6365715
GENERAL INFORMATION:
APPLICANT: ARLETH, ANTHONY J.
APPLICANT: WILLETTTE, ROBERT N.
APPLICANT: ELSHOURBAGY, NABIL A.
APPLICANT: LI, XIAOTONG
TITLE OF INVENTION: HUMAN CARDIAC/BRAIN TOLLDOID-LIKE PROTEIN
FILE REFERENCE: ATG-50038-D1
CURRENT APPLICATION NUMBER: US/09/432,473
CURRENT FILING DATE: 1999-11-01
EARLIER APPLICATION NUMBER: 08/991,408
EARLIER FILING DATE: 1997-12-16
EARLIER APPLICATION NUMBER: 60/034,471
EARLIER FILING DATE: 1997-01-02
NUMBER OF SEQ ID NOS: 4
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 4
LENGTH: 591
TYPE: PRT
ORGANISM: HOMO SAPIENS
US-09-432-473-4

Query Match 8.7%; Score 172.5; DB 4; Length 591;
Best Local Similarity 36.7%; Pred. No. 4.6e-09;
Matches 40; Conservative 25; Mismatches 39; Indels 5; Gaps 4;

QY 59 KGNQVQSPRPSPYPRNLLITRLHSQENTRIQLVDFDNOFGLEAENDICRDEYVED 118
Db 202 KLNQGITTPGMPREYEPNKNKCIQVAPTOYRISVFE--FEELE--GNEVCKYDYEIWS 258
QY 119 ISETSTIRGRCMGHKEVPPRIKSRNQIKITFKSDDYFAKPGFKIYY 167
Db 259 GLSESKLHGKFCG-AEVEVITSQYNNMVEFEKSDN-TVSKKGKFAHF 305

RESULT 15
US-08-866-650-5
Sequence 5, Application US/08866650
Patent No. 5938321
GENERAL INFORMATION:
APPLICANT: Greenspan, Daniel S
APPLICANT: Takahara, Kazuhiko
APPLICANT: Hoffman, Guy G
TITLE OF INVENTION: Mammalian Tollold-Like Protein
NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: Quarles & Brady
STREET: 1 South Plinckney Street
CITY: Madison
STATE: WI
COUNTRY: US
ZIP: 53703
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30

GenCore version 5.1.6
Copyright (c) 1993 - 2003 CompuGen Ltd.

OM protein - protein search, using sw model

Run on: June 12, 2003, 15:32:31 ; Search time 22 Seconds

(Without alignments)
1736.314 Million cell updates/sec

Title: US-09-662-783-2

Perfect score: 1994

Sequence: 1 MHRILFYVTLICANFCSCRD.....DIQLDHERCDICSSNPR 370

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 392085 seqs, 103240269 residues

Total number of hits satisfying chosen parameters: 392085

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Published_Applications_AA:*
1: /cgn2_6/ptodata/2/pubppaa/US08_NEW_PUB pep:*
2: /cgn2_6/ptodata/2/pubppaa/PCT_NEW_PUB pep:*
3: /cgn2_6/ptodata/2/pubppaa/US06_PUBCOMB pep:*
4: /cgn2_6/ptodata/2/pubppaa/US07_NEW_PUB pep:*
5: /cgn2_6/ptodata/2/pubppaa/US07_PUBCOMB pep:*
6: /cgn2_6/ptodata/2/pubppaa/PCTUS_PUBCOMB pep:*
7: /cgn2_6/ptodata/2/pubppaa/US08_PUBCOMB pep:*
8: /cgn2_6/ptodata/2/pubppaa/US09_NEW_PUB pep:*
9: /cgn2_6/ptodata/2/pubppaa/US09_PUBCOMB pep:*
10: /cgn2_6/ptodata/2/pubppaa/US10_NEW_PUB pep:*
11: /cgn2_6/ptodata/2/pubppaa/US10_PUBCOMB pep:*
12: /cgn2_6/ptodata/2/pubppaa/US60_NEW_PUB pep:*
13: /cgn2_6/ptodata/2/pubppaa/US60_PUBCOMB pep:*
14: /cgn2_6/ptodata/2/pubppaa/US60_PUBCOMB pep:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	1994	100.0	370	9 US-10-086-623-8	Sequence 8, Appli
2	1994	100.0	370	9 US-10-139-583-37	Sequence 37, Appli
3	1994	100.0	370	9 US-10-039-847A-2	Sequence 2, Appli
4	1994	100.0	370	9 US-10-260-539-8	Sequence 8, Appli
5	1994	100.0	370	9 US-10-258-557-2	Sequence 5, Appli
6	1994	100.0	370	10 US-09-823-033-5	Sequence 2, Appli
7	1994	100.0	370	10 US-09-808-972-2	Sequence 5, Appli
8	1994	100.0	370	10 US-09-915-582-56	Sequence 2, Appli
9	1994	100.0	370	10 US-09-915-582-74	Sequence 56, Appli
10	1988	99.7	364	9 US-10-028-072-186	Sequence 74, Appli
11	1949	97.7	364	9 US-10-121-049-186	Sequence 186, App
12	1949	97.7	364	9 US-10-123-904-186	Sequence 186, App
13	1949	97.7	364	9 US-10-140-470-186	Sequence 186, App
14	1949	97.7	364	9 US-10-175-746-186	Sequence 186, App
15	1949	97.7	364	9 US-10-176-918-186	Sequence 186, App
16	1949	97.7	364	9 US-10-176-921-186	Sequence 186, App
17	1949	97.7	364	9 US-10-137-865-186	Sequence 186, App
18	1949	97.7	364	9 US-10-140-474-186	Sequence 186, App
19	1949	97.7	364	9 US-10-140-474-186	Sequence 186, App

20	1949	97.7	364	9 US-10-142-431-186	Sequence 186, App
21	1949	97.7	364	9 US-10-143-114-186	Sequence 186, App
22	1949	97.7	364	9 US-10-140-002-186	Sequence 186, App
23	1949	97.7	364	9 US-10-142-419-186	Sequence 186, App
24	1949	97.7	364	9 US-10-123-262-186	Sequence 186, App
25	1949	97.7	364	9 US-10-142-423-186	Sequence 186, App
26	1949	97.7	364	9 US-10-121-050-186	Sequence 186, App
27	1949	97.7	364	9 US-10-141-755-186	Sequence 186, App
28	1949	97.7	364	9 US-10-143-032-186	Sequence 186, App
29	1949	97.7	364	9 US-10-123-108-186	Sequence 186, App
30	1949	97.7	364	9 US-10-123-236-186	Sequence 186, App
31	1949	97.7	364	9 US-10-123-261-186	Sequence 186, App
32	1949	97.7	364	9 US-10-140-921-186	Sequence 186, App
33	1949	97.7	364	9 US-10-140-928-186	Sequence 186, App
34	1949	97.7	364	9 US-10-121-045-186	Sequence 186, App
35	1949	97.7	364	9 US-10-123-292-186	Sequence 186, App
36	1949	97.7	364	9 US-10-123-903-186	Sequence 186, App
37	1949	97.7	364	9 US-10-124-819-186	Sequence 186, App
38	1949	97.7	364	9 US-10-124-822-186	Sequence 186, App
39	1949	97.7	364	9 US-10-140-925-186	Sequence 186, App
40	1949	97.7	364	9 US-10-160-498-186	Sequence 186, App
41	1949	97.7	364	9 US-10-121-041-186	Sequence 186, App
42	1949	97.7	364	9 US-10-121-043-186	Sequence 186, App
43	1949	97.7	364	9 US-10-121-047-186	Sequence 186, App
44	1949	97.7	364	9 US-10-123-215-186	Sequence 186, App
45	1949	97.7	364	9 US-10-123-902-186	Sequence 186, App

ALIGNMENTS

RESULT 1
US-10-086-623-8
Sequence 8, Application US/10086623
Patent No. US20020164710A1
GENERAL INFORMATION:
APPLICANT: ERIKSSON, Ulf
APPLICANT: AASE, Karin
APPLICANT: LI, Xuri
APPLICANT: FONTEN, Annica
APPLICANT: UTELLA, Marko
APPLICANT: ALITALO, Karl
APPLICANT: OESTMAN, Arne
APPLICANT: HELDIN, Carl-Henrik
TITLE OF INVENTION: PLATELET DERIVED GROWTH FACTOR D, DNA CODING THEREFOR AND USES
FILE REFERENCE: 1064/44833C2
CURRENT APPLICATION NUMBER: US/10/086, 623
CURRENT FILING DATE: 2000-03-04
PRIOR APPLICATION NUMBER: US 60/107, 852
PRIOR FILING DATE: 1998-11-10
PRIOR APPLICATION NUMBER: US 60/113, 997
PRIOR FILING DATE: 1998-12-28
PRIOR APPLICATION NUMBER: US 60/150, 604
PRIOR FILING DATE: 1999-08-26
PRIOR APPLICATION NUMBER: US 60/157, 108
PRIOR FILING DATE: 1999-10-04
PRIOR APPLICATION NUMBER: US 60/157, 756
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: US 09/438, 046
PRIOR FILING DATE: 1999-11-10
PRIOR APPLICATION NUMBER: US 09/691, 200
PRIOR FILING DATE: 2000-10-19
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PatentIn version 3.1
SEQ ID NO 8
LENGTH: 370
TYPE: PRT
ORGANISM: Homo sapiens
US-10-086-623-8
Query Match 100.0%; Score 1994; DB 9; Length 370;
Best Local Similarity 100.0%; Pred. No. 3e-161;
Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MRLIFVYLLICANFCSGCRDTSATPOSASIKALRNANLRDESNHLLTDLYRDETIQVKG 60
1 MRLIFVYLLICANFCSGCRDTSATPOSASIKALRNANLRDESNHLLTDLYRDETIQVKG 60
Db 61 NGVOSPRFPNSYPRNLLTWRLHSEENRIQVFNQGLAEANDICRYDEVEDIS 120
61 NGVOSPRFPNSYPRNLLTWRLHSEENRIQVFNQGLAEANDICRYDEVEDIS 120
QY 121 ESTIIRGRWCGHKEVPPRIKSRTOIKITFKSDDYFVAKPGFKIYSLLEDFOPAAASE 180
121 ESTIIRGRWCGHKEVPPRIKSRTOIKITFKSDDYFVAKPGFKIYSLLEDFOPAAASE 180
Db 121 ESTIIRGRWCGHKEVPPRIKSRTOIKITFKSDDYFVAKPGFKIYSLLEDFOPAAASE 180
121 ESTIIRGRWCGHKEVPPRIKSRTOIKITFKSDDYFVAKPGFKIYSLLEDFOPAAASE 180
QY 181 TMESTYSSISGVSYNSPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESWQEDLENNY 240
181 TMESTYSSISGVSYNSPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESWQEDLENNY 240
Db 241 LDPFRGRSYHDKRSKVDLDRLNDADAKRSCPTPRNYSVIREELKIANVFFPRCLVQ 300
241 LDPFRGRSYHDKRSKVDLDRLNDADAKRSCPTPRNYSVIREELKIANVFFPRCLVQ 300
QY 301 RCGNGCGGVNMRSCCTCNSGKTVKKYHEVLOFEPGHIKRGAKTMALVDIOLDHHERC 360
301 RCGNGCGGVNMRSCCTCNSGKTVKKYHEVLOFEPGHIKRGAKTMALVDIOLDHHERC 360
Db 301 RCGNGCGGVNMRSCCTCNSGKTVKKYHEVLOFEPGHIKRGAKTMALVDIOLDHHERC 360
301 RCGNGCGGVNMRSCCTCNSGKTVKKYHEVLOFEPGHIKRGAKTMALVDIOLDHHERC 360
QY 361 DCICSSRPPR 370
361 DCICSSRPPR 370
Db 361 DCICSSRPPR 370
361 DCICSSRPPR 370

RESULT 2
US-10-139-583-37
Sequence 37, Application US/10139583
Patent No. US20020177193A1
GENERAL INFORMATION:
APPLICANT: Gao, Zeren
APPLICANT: Hart, Charles E.
APPLICANT: Piddington, Christopher S.
APPLICANT: Sheppard, Paul O.
APPLICANT: Shoemaker, Kimberly E.
APPLICANT: Gilbertson, Debra G.
APPLICANT: West, James W.
TITLE OF INVENTION: GROWTH FACTOR HOMOLOG ZVEG33
FILE REFERENCE: 98-60
CURRENT APPLICATION NUMBER: US/10/139,583
CURRENT FILING DATE: 2002-05-02
PRIOR APPLICATION NUMBER: 09/457,066
PRIOR FILING DATE: 1999-12-07
NUMBER OF SEQ ID NOS: 50
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 37
LENGTH: 370
TYPE: PRN
ORGANISM: Homo sapiens
US-10-139-583-37

Query Match 100.0%; Score 1994; DB 9; Length 370;
Best Local Similarity 100.0%; Pred. No. 3e-161;
Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MRLIFVYLLICANFCSGCRDTSATPOSASIKALRNANLRDESNHLLTDLYRDETIQVKG 60
1 MRLIFVYLLICANFCSGCRDTSATPOSASIKALRNANLRDESNHLLTDLYRDETIQVKG 60
Db 61 NGVOSPRFPNSYPRNLLTWRLHSEENRIQVFNQGLAEANDICRYDEVEDIS 120
61 NGVOSPRFPNSYPRNLLTWRLHSEENRIQVFNQGLAEANDICRYDEVEDIS 120
QY 121 ESTIIRGRWCGHKEVPPRIKSRTOIKITFKSDDYFVAKPGFKIYSLLEDFOPAAASE 180
121 ESTIIRGRWCGHKEVPPRIKSRTOIKITFKSDDYFVAKPGFKIYSLLEDFOPAAASE 180
Db 121 ESTIIRGRWCGHKEVPPRIKSRTOIKITFKSDDYFVAKPGFKIYSLLEDFOPAAASE 180
121 ESTIIRGRWCGHKEVPPRIKSRTOIKITFKSDDYFVAKPGFKIYSLLEDFOPAAASE 180
QY 181 TMESTYSSISGVSYNSPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESWQEDLENNY 240
181 TMESTYSSISGVSYNSPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESWQEDLENNY 240

Db 181 TMESTYSSISGVSYNSPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESWQEDLENNY 240
181 TMESTYSSISGVSYNSPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESWQEDLENNY 240
QY 241 LDPFRGRSYHDKRSKVDLDRLNDADAKRSCPTPRNYSVIREELKIANVFFPRCLVQ 300
241 LDPFRGRSYHDKRSKVDLDRLNDADAKRSCPTPRNYSVIREELKIANVFFPRCLVQ 300
Db 241 LDPFRGRSYHDKRSKVDLDRLNDADAKRSCPTPRNYSVIREELKIANVFFPRCLVQ 300
241 LDPFRGRSYHDKRSKVDLDRLNDADAKRSCPTPRNYSVIREELKIANVFFPRCLVQ 300
QY 301 RCGNGCGGVNMRSCCTCNSGKTVKKYHEVLOFEPGHIKRGAKTMALVDIOLDHHERC 360
301 RCGNGCGGVNMRSCCTCNSGKTVKKYHEVLOFEPGHIKRGAKTMALVDIOLDHHERC 360
Db 301 RCGNGCGGVNMRSCCTCNSGKTVKKYHEVLOFEPGHIKRGAKTMALVDIOLDHHERC 360
301 RCGNGCGGVNMRSCCTCNSGKTVKKYHEVLOFEPGHIKRGAKTMALVDIOLDHHERC 360
QY 361 DCICSSRPPR 370
361 DCICSSRPPR 370
Db 361 DCICSSRPPR 370
361 DCICSSRPPR 370

RESULT 3
US-10-039-847A-2
Sequence 2, Application US/10039847A
Publication No. US20020183273A1
GENERAL INFORMATION:
APPLICANT: Hart, Charles E.
APPLICANT: Popouzis, Stavros
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR IMPROVING
FILE REFERENCE: 00-100
CURRENT APPLICATION NUMBER: US/10/039,847A
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: US 60/244,479
PRIOR FILING DATE: 2000-10-30
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 2
LENGTH: 370
TYPE: PRN
ORGANISM: Homo sapiens
US-10-039-847A-2

Query Match 100.0%; Score 1994; DB 9; Length 370;
Best Local Similarity 100.0%; Pred. No. 3e-161;
Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MRLIFVYLLICANFCSGCRDTSATPOSASIKALRNANLRDESNHLLTDLYRDETIQVKG 60
1 MRLIFVYLLICANFCSGCRDTSATPOSASIKALRNANLRDESNHLLTDLYRDETIQVKG 60
Db 61 NGVOSPRFPNSYPRNLLTWRLHSEENRIQVFNQGLAEANDICRYDEVEDIS 120
61 NGVOSPRFPNSYPRNLLTWRLHSEENRIQVFNQGLAEANDICRYDEVEDIS 120
QY 121 ESTIIRGRWCGHKEVPPRIKSRTOIKITFKSDDYFVAKPGFKIYSLLEDFOPAAASE 180
121 ESTIIRGRWCGHKEVPPRIKSRTOIKITFKSDDYFVAKPGFKIYSLLEDFOPAAASE 180
Db 121 ESTIIRGRWCGHKEVPPRIKSRTOIKITFKSDDYFVAKPGFKIYSLLEDFOPAAASE 180
121 ESTIIRGRWCGHKEVPPRIKSRTOIKITFKSDDYFVAKPGFKIYSLLEDFOPAAASE 180
QY 181 TMESTYSSISGVSYNSPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESWQEDLENNY 240
181 TMESTYSSISGVSYNSPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESWQEDLENNY 240
Db 241 LDPFRGRSYHDKRSKVDLDRLNDADAKRSCPTPRNYSVIREELKIANVFFPRCLVQ 300
241 LDPFRGRSYHDKRSKVDLDRLNDADAKRSCPTPRNYSVIREELKIANVFFPRCLVQ 300
QY 301 RCGNGCGGVNMRSCCTCNSGKTVKKYHEVLOFEPGHIKRGAKTMALVDIOLDHHERC 360
301 RCGNGCGGVNMRSCCTCNSGKTVKKYHEVLOFEPGHIKRGAKTMALVDIOLDHHERC 360
Db 301 RCGNGCGGVNMRSCCTCNSGKTVKKYHEVLOFEPGHIKRGAKTMALVDIOLDHHERC 360
301 RCGNGCGGVNMRSCCTCNSGKTVKKYHEVLOFEPGHIKRGAKTMALVDIOLDHHERC 360
QY 361 DCICSSRPPR 370
361 DCICSSRPPR 370
Db 361 DCICSSRPPR 370
361 DCICSSRPPR 370

RESULT 4
US-10-260-539-8

```
Sequence 8, Application US/10260539
: Publication No. US20030073637A1
: GENERAL INFORMATION:
: APPLICANT: ERIKSSON, Ulf
: APPLICANT: AASE, Karin
: APPLICANT: LI, Xuri
: APPLICANT: PONTEN, Annica
: APPLICANT: UUTELA, Marko
: APPLICANT: ALITALO, Karl
: APPLICANT: OESTMAN, Arne
: APPLICANT: HELDIN, Carl-Henrik
: TITLE OF INVENTION: PLATELET DERIVED GROWTH FACTOR D, DNA CODING THEREFOR AND USES TH
: FILE REFERENCE: 1064/44833C2
: CURRENT APPLICATION NUMBER: US/10/260,539
: CURRENT FILING DATE: 2002-10-01
: PRIOR APPLICATION NUMBER: US/10/086,623
: PRIOR FILING DATE: 2000-03-04
: PRIOR APPLICATION NUMBER: US 60/107,852
: PRIOR FILING DATE: 1998-11-10
: PRIOR APPLICATION NUMBER: US 60/113,997
: PRIOR FILING DATE: 1998-12-28
: PRIOR APPLICATION NUMBER: US 60/150,604
: PRIOR FILING DATE: 1999-08-26
: PRIOR APPLICATION NUMBER: US 60/157,108
: PRIOR FILING DATE: 1999-10-04
: PRIOR APPLICATION NUMBER: US 60/157,756
: PRIOR FILING DATE: 1999-10-05
: PRIOR APPLICATION NUMBER: US 09/438,046
: PRIOR FILING DATE: 1999-11-10
: PRIOR APPLICATION NUMBER: US 09/691,200
: PRIOR FILING DATE: 2000-10-19
: NUMBER OF SEQ ID NOS: 42
: SOFTWARE: PatentIn version 3.1
: SEQ ID NO 8
: LENGTH: 370
: TYPE: PRT
: ORGANISM: Homo sapiens
: US-10-260-539-8

Query Match          100.0%; Score 1994; DB 9; Length 370;
Best Local Similarity 100.0%; Pred. No. 3e-161;
Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MHRLEFYTLICANFCSCRODTSATPOSASIKALNNANLRDESNHLDLYRDETIQYKG 60
DB 1 MHRLEFYTLICANFCSCRODTSATPOSASIKALNNANLRDESNHLDLYRDETIQYKG 60
QY 61 NGYQSPREFPNSYRNLLTWRLHSEENTRIQLVFNQFGLFEAENDICRYDVEVEDIS 120
DB 61 NGYQSPREFPNSYRNLLTWRLHSEENTRIQLVFNQFGLFEAENDICRYDVEVEDIS 120
QY 121 ESTIIRGRWCGRHKEVPRIKSRINOIKITFKSDYFAKGFYIYSLDFQPAASE 180
DB 121 ESTIIRGRWCGRHKEVPRIKSRINOIKITFKSDYFAKGFYIYSLDFQPAASE 180
QY 121 TMSVTSISGVSNSPSVTPDLADLADKTAEPFTVDLKYFNPESMOEDLEMY 240
DB 121 TMSVTSISGVSNSPSVTPDLADLADKTAEPFTVDLKYFNPESMOEDLEMY 240
QY 181 TMSVTSISGVSNSPSVTPDLADLADKTAEPFTVDLKYFNPESMOEDLEMY 240
DB 181 TMSVTSISGVSNSPSVTPDLADLADKTAEPFTVDLKYFNPESMOEDLEMY 240
QY 241 LDPFRYGRSYHDKRSKYDLRLNDADKARYSCPTPRNSVNIREEIKLANVFFPCLLVQ 300
DB 241 LDPFRYGRSYHDKRSKYDLRLNDADKARYSCPTPRNSVNIREEIKLANVFFPCLLVQ 300
QY 301 RCGNCGCGTVNMRSCTCNSGKTYKKYHEVLQEPFGHKKRGRAKTMAVLDIQLDHERC 360
DB 301 RCGNCGCGTVNMRSCTCNSGKTYKKYHEVLQEPFGHKKRGRAKTMAVLDIQLDHERC 360
QY 301 RCGNCGCGTVNMRSCTCNSGKTYKKYHEVLQEPFGHKKRGRAKTMAVLDIQLDHERC 360
DB 301 RCGNCGCGTVNMRSCTCNSGKTYKKYHEVLQEPFGHKKRGRAKTMAVLDIQLDHERC 360
QY 361 DCICSSRPR 370
DB 361 DCICSSRPR 370

RESULT 5
```

```
US-10-264-361-5
: Sequence 5, Application US/10264361
: Publication No. US20030087870A1
: GENERAL INFORMATION:
: APPLICANT: GILBERTSON, Debra G.
: TITLE OF INVENTION: METHOD OF TREATING FIBROSIS
: FILE REFERENCE: 00-53
: CURRENT APPLICATION NUMBER: US/10/264,361
: CURRENT FILING DATE: 2002-10-03
: PRIOR APPLICATION NUMBER: US/09/695,121
: PRIOR FILING DATE: 2000-10-23
: NUMBER OF SEQ ID NOS: 18
: SOFTWARE: FastSeq for Windows Version 3.0
: SEQ ID NO 5
: LENGTH: 370
: TYPE: PRT
: ORGANISM: Homo sapiens
: US-10-264-361-5

Query Match          100.0%; Score 1994; DB 9; Length 370;
Best Local Similarity 100.0%; Pred. No. 3e-161;
Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MHRLEFYTLICANFCSCRODTSATPOSASIKALNNANLRDESNHLDLYRDETIQYKG 60
DB 1 MHRLEFYTLICANFCSCRODTSATPOSASIKALNNANLRDESNHLDLYRDETIQYKG 60
QY 61 NGYQSPREFPNSYRNLLTWRLHSEENTRIQLVFNQFGLFEAENDICRYDVEVEDIS 120
DB 61 NGYQSPREFPNSYRNLLTWRLHSEENTRIQLVFNQFGLFEAENDICRYDVEVEDIS 120
QY 121 ESTIIRGRWCGRHKEVPRIKSRINOIKITFKSDYFAKGFYIYSLDFQPAASE 180
DB 121 ESTIIRGRWCGRHKEVPRIKSRINOIKITFKSDYFAKGFYIYSLDFQPAASE 180
QY 121 TMSVTSISGVSNSPSVTPDLADLADKTAEPFTVDLKYFNPESMOEDLEMY 240
DB 121 TMSVTSISGVSNSPSVTPDLADLADKTAEPFTVDLKYFNPESMOEDLEMY 240
QY 241 LDPFRYGRSYHDKRSKYDLRLNDADKARYSCPTPRNSVNIREEIKLANVFFPCLLVQ 300
DB 241 LDPFRYGRSYHDKRSKYDLRLNDADKARYSCPTPRNSVNIREEIKLANVFFPCLLVQ 300
QY 301 RCGNCGCGTVNMRSCTCNSGKTYKKYHEVLQEPFGHKKRGRAKTMAVLDIQLDHERC 360
DB 301 RCGNCGCGTVNMRSCTCNSGKTYKKYHEVLQEPFGHKKRGRAKTMAVLDIQLDHERC 360
QY 361 DCICSSRPR 370
DB 361 DCICSSRPR 370

RESULT 6
US-10-258-557-2
: Sequence 2, Application US/10258557
: Publication No. US20030100502A1
: GENERAL INFORMATION:
: APPLICANT: BEALS, John
: APPLICANT: GONZALEZ-DEMILLT, Patricia
: APPLICANT: HAMMOND, Lisa
: APPLICANT: LU, Jitong
: APPLICANT: NA, Songqing
: APPLICANT: SU, Eric
: APPLICANT: WITCHER, Derrick
: TITLE OF INVENTION: TREATING MUSCULOSKELETAL DISORDERS USING LP85 AND ANALOGS THERE
: FILE REFERENCE: X-14392M
: CURRENT APPLICATION NUMBER: US/10/258,557
: CURRENT FILING DATE: 2002-10-23
: NUMBER OF SEQ ID NOS: 6
: SOFTWARE: PatentIn version 3.0
: SEQ ID NO 2
: LENGTH: 370
: TYPE: PRT
```

ORGANISM: Homo sapiens
US-10-258-557-2

Query Match 100.0%; Score 1994; DB 9; Length 370;
Best Local Similarity 100.0%; Pred. No. 3e-161;
Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MHRLLFYVTLICANFCSCRDTSATPOSASIKALRNANLRDESNHLLTDLRRDETIOVKG 60
DB 1 MHRLLFYVTLICANFCSCRDTSATPOSASIKALRNANLRDESNHLLTDLRRDETIOVKG 60
QY 61 NGVOSPRPNSTPRNLLTLWRHLSOENRIQLVFNQGLBEAENDICRYDVEVEDIS 120
DB 61 NGVOSPRPNSTPRNLLTLWRHLSOENRIQLVFNQGLBEAENDICRYDVEVEDIS 120
QY 121 ESTIIRGRCGHEKVEPPRIKSTNOIKITFKSDDYFVAKPGFKIYSLLEDPQPAASE 180
DB 121 ESTIIRGRCGHEKVEPPRIKSTNOIKITFKSDDYFVAKPGFKIYSLLEDPQPAASE 180
QY 181 TMWESVTSSISGVSNPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESMOEDLENNY 240
DB 181 TMWESVTSSISGVSNPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESMOEDLENNY 240
QY 241 LDPFRGRSYHDKRSKVDLRLNDAKRISCTPRNYSVINIRELKLAVVFFPRLVQ 300
DB 241 LDPFRGRSYHDKRSKVDLRLNDAKRISCTPRNYSVINIRELKLAVVFFPRLVQ 300
QY 301 RCGNGCGCTVMNRSCCTNSGKTGKYYHEVLOPEPGHKKRGRAKTALVDIOLDHHERC 360
DB 301 RCGNGCGCTVMNRSCCTNSGKTGKYYHEVLOPEPGHKKRGRAKTALVDIOLDHHERC 360
QY 361 DCICSSRPPR 370
DB 361 DCICSSRPPR 370

RESULT 7

US-09-823-033-5
Sequence 5, Application US/09823033
Patent No. US20020004225A1
GENERAL INFORMATION:
APPLICANT: Hart, Charles E.
TITLE OF INVENTION: METHODS FOR PROMOTING GROWTH OF BONE,
FILE REFERENCE: 00-12
CURRENT APPLICATION NUMBER: US/09/823,033
CURRENT FILING DATE: 2001-03-29
NUMBER OF SEQ ID NOS: 5
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 5
LENGTH: 370
TYPE: PRN
ORGANISM: Homo sapiens
US-09-823-033-5

Query Match 100.0%; Score 1994; DB 10; Length 370;
Best Local Similarity 100.0%; Pred. No. 3e-161;
Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MHRLLFYVTLICANFCSCRDTSATPOSASIKALRNANLRDESNHLLTDLRRDETIOVKG 60
DB 1 MHRLLFYVTLICANFCSCRDTSATPOSASIKALRNANLRDESNHLLTDLRRDETIOVKG 60
QY 61 NGVOSPRPNSTPRNLLTLWRHLSOENRIQLVFNQGLBEAENDICRYDVEVEDIS 120
DB 61 NGVOSPRPNSTPRNLLTLWRHLSOENRIQLVFNQGLBEAENDICRYDVEVEDIS 120
QY 121 ESTIIRGRCGHEKVEPPRIKSTNOIKITFKSDDYFVAKPGFKIYSLLEDPQPAASE 180
DB 121 ESTIIRGRCGHEKVEPPRIKSTNOIKITFKSDDYFVAKPGFKIYSLLEDPQPAASE 180
QY 181 TMWESVTSSISGVSNPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESMOEDLENNY 240
DB 181 TMWESVTSSISGVSNPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESMOEDLENNY 240

|||||
DB 181 TMWESVTSSISGVSNPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESMOEDLENNY 240

QY 241 LDPFRGRSYHDKRSKVDLRLNDAKRISCTPRNYSVINIRELKLAVVFFPRLVQ 300
DB 241 LDPFRGRSYHDKRSKVDLRLNDAKRISCTPRNYSVINIRELKLAVVFFPRLVQ 300
QY 301 RCGNGCGCTVMNRSCCTNSGKTGKYYHEVLOPEPGHKKRGRAKTALVDIOLDHHERC 360
DB 301 RCGNGCGCTVMNRSCCTNSGKTGKYYHEVLOPEPGHKKRGRAKTALVDIOLDHHERC 360
QY 361 DCICSSRPPR 370
DB 361 DCICSSRPPR 370

RESULT 8

US-09-808-972-2
Sequence 2, Application US/09808972
Patent No. US20020064832A1
GENERAL INFORMATION:
APPLICANT: Hart, Charles E.
APPLICANT: Topouzis, Stavros
TITLE OF INVENTION: METHOD OF TREATING FIBROPROLIFERATIVE
FILE REFERENCE: 00-79
CURRENT APPLICATION NUMBER: US/09/808,972
CURRENT FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: US 60/235,295
PRIOR FILING DATE: 2000-09-26
PRIOR APPLICATION NUMBER: US 09/564,595
PRIOR FILING DATE: 2000-05-03
PRIOR APPLICATION NUMBER: US 60/180,169
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/164,463
PRIOR FILING DATE: 1999-11-10
PRIOR APPLICATION NUMBER: US 60/132,250
PRIOR FILING DATE: 1999-05-03
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 2
LENGTH: 370
TYPE: PRN
ORGANISM: Homo sapiens
US-09-808-972-2

Query Match 100.0%; Score 1994; DB 10; Length 370;
Best Local Similarity 100.0%; Pred. No. 3e-161;
Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MHRLLFYVTLICANFCSCRDTSATPOSASIKALRNANLRDESNHLLTDLRRDETIOVKG 60
DB 1 MHRLLFYVTLICANFCSCRDTSATPOSASIKALRNANLRDESNHLLTDLRRDETIOVKG 60
QY 61 NGVOSPRPNSTPRNLLTLWRHLSOENRIQLVFNQGLBEAENDICRYDVEVEDIS 120
DB 61 NGVOSPRPNSTPRNLLTLWRHLSOENRIQLVFNQGLBEAENDICRYDVEVEDIS 120
QY 121 ESTIIRGRCGHEKVEPPRIKSTNOIKITFKSDDYFVAKPGFKIYSLLEDPQPAASE 180
DB 121 ESTIIRGRCGHEKVEPPRIKSTNOIKITFKSDDYFVAKPGFKIYSLLEDPQPAASE 180
QY 181 TMWESVTSSISGVSNPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESMOEDLENNY 240
DB 181 TMWESVTSSISGVSNPSVTDPTLLADALDKKIAEFDVEDLLKTFNPESMOEDLENNY 240
QY 241 LDPFRGRSYHDKRSKVDLRLNDAKRISCTPRNYSVINIRELKLAVVFFPRLVQ 300
DB 241 LDPFRGRSYHDKRSKVDLRLNDAKRISCTPRNYSVINIRELKLAVVFFPRLVQ 300
QY 301 RCGNGCGCTVMNRSCCTNSGKTGKYYHEVLOPEPGHKKRGRAKTALVDIOLDHHERC 360
DB 301 RCGNGCGCTVMNRSCCTNSGKTGKYYHEVLOPEPGHKKRGRAKTALVDIOLDHHERC 360

Db 301 RCGNCGGTYNMNRCTCNSGKTVKKYHEVLQEPFGHIKRRGAKTALVDIQDHHERC 360
Qy 361 DCICSSRPPR 370
Db 361 DCICSSRPPR 370

RESULT 9

US-09-915-582-56
; Sequence 56, Application US/09915582
; Patent No. US20020120103A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: 17 Human Secreted Proteins
; FILE REFERENCE: PS723P1
; CURRENT APPLICATION NUMBER: US/09/915,582
; CURRENT FILING DATE: 2001-07-27
; PRIOR APPLICATION NUMBER: PCT/US01/01431
; PRIOR FILING DATE: 2001-01-17
; PRIOR APPLICATION NUMBER: 60/179,065
; PRIOR FILING DATE: 2000-01-31
; PRIOR APPLICATION NUMBER: 60/180,628
; PRIOR FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: 60/231,968
; PRIOR FILING DATE: 2000-09-12
; NUMBER OF SEQ ID NOS: 97
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 56
; LENGTH: 370
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-915-582-56

Query Match 100.0%; Score 1994; DB 10; Length 370;
Best Local Similarity 100.0%; Pred. No. 3e-161;
Matches 370; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MHRLLFYTLICANFCSCROTSATPOSASIKALRNANLRDESHNLDLYRDETIQYKG 60
Db 1 MHRLLFYTLICANFCSCROTSATPOSASIKALRNANLRDESHNLDLYRDETIQYKG 60
Qy 61 NGYQSPRPSPSYRNLLTWRLHSEENTRIQLVFNQFGLAEANDICRYDFVEVEDIS 120
Db 61 NGYQSPRPSPSYRNLLTWRLHSEENTRIQLVFNQFGLAEANDICRYDFVEVEDIS 120
Qy 121 ETSITLGRMCGHKEVPRIKSRNQIKITFKSDDYFAKGFITYSLDEFPQAAASE 180
Db 121 ETSITLGRMCGHKEVPRIKSRNQIKITFKSDDYFAKGFITYSLDEFPQAAASE 180
Qy 181 TNMESVTSISGVSNPSVTDPLIADALDKKTAEPDVTEDLAKYFNPESMODLEMY 240
Db 181 TNMESVTSISGVSNPSVTDPLIADALDKKTAEPDVTEDLAKYFNPESMODLEMY 240
Qy 241 LDTPRYGRSYHDKRSKYVDLRLNDADKRYSCPTPRNYSVNIREEKLANVVEFPFCLLVQ 300
Db 241 LDTPRYGRSYHDKRSKYVDLRLNDADKRYSCPTPRNYSVNIREEKLANVVEFPFCLLVQ 300
Qy 301 RCGNCGGTYNMNRCTCNSGKTVKKYHEVLQEPFGHIKRRGAKTALVDIQDHHERC 360
Db 301 RCGNCGGTYNMNRCTCNSGKTVKKYHEVLQEPFGHIKRRGAKTALVDIQDHHERC 360
Qy 361 DCICSSRPPR 370
Db 361 DCICSSRPPR 370

RESULT 10

US-09-915-582-74
; Sequence 74, Application US/09915582
; Patent No. US20020120103A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: 17 Human Secreted Proteins

; FILE REFERENCE: PS723P1
; CURRENT APPLICATION NUMBER: US/09/915,582
; CURRENT FILING DATE: 2001-07-27
; PRIOR APPLICATION NUMBER: PCT/US01/01431
; PRIOR FILING DATE: 2001-01-17
; PRIOR APPLICATION NUMBER: 60/179,065
; PRIOR FILING DATE: 2000-01-31
; PRIOR APPLICATION NUMBER: 60/180,628
; PRIOR FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: 60/231,968
; PRIOR FILING DATE: 2000-09-12
; NUMBER OF SEQ ID NOS: 97
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 74
; LENGTH: 370
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (216)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-09-915-582-74

Query Match 99.7%; Score 1988; DB 10; Length 370;
Best Local Similarity 99.7%; Pred. No. 9.6e-161;
Matches 369; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MHRLLFYTLICANFCSCROTSATPOSASIKALRNANLRDESHNLDLYRDETIQYKG 60
Db 1 MHRLLFYTLICANFCSCROTSATPOSASIKALRNANLRDESHNLDLYRDETIQYKG 60
Qy 61 NGYQSPRPSPSYRNLLTWRLHSEENTRIQLVFNQFGLAEANDICRYDFVEVEDIS 120
Db 61 NGYQSPRPSPSYRNLLTWRLHSEENTRIQLVFNQFGLAEANDICRYDFVEVEDIS 120
Qy 121 ETSITLGRMCGHKEVPRIKSRNQIKITFKSDDYFAKGFITYSLDEFPQAAASE 180
Db 121 ETSITLGRMCGHKEVPRIKSRNQIKITFKSDDYFAKGFITYSLDEFPQAAASE 180
Qy 181 TNMESVTSISGVSNPSVTDPLIADALDKKTAEPDVTEDLAKYFNPESMODLEMY 240
Db 181 TNMESVTSISGVSNPSVTDPLIADALDKKTAEPDVTEDLAKYFNPESMODLEMY 240
Qy 241 LDTPRYGRSYHDKRSKYVDLRLNDADKRYSCPTPRNYSVNIREEKLANVVEFPFCLLVQ 300
Db 241 LDTPRYGRSYHDKRSKYVDLRLNDADKRYSCPTPRNYSVNIREEKLANVVEFPFCLLVQ 300
Qy 301 RCGNCGGTYNMNRCTCNSGKTVKKYHEVLQEPFGHIKRRGAKTALVDIQDHHERC 360
Db 301 RCGNCGGTYNMNRCTCNSGKTVKKYHEVLQEPFGHIKRRGAKTALVDIQDHHERC 360
Qy 361 DCICSSRPPR 370
Db 361 DCICSSRPPR 370

RESULT 11

US-10-028-072-186
; Sequence 186, Application US/10028072
; Publication No. US20030004311A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
; APPLICANT: Beresini, Maureen
; APPLICANT: DeForge, Laura
; APPLICANT: Desnoyers, Luc
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Sherwood, Steven
; APPLICANT: Smith, Victoria

APPLICANT: Stewart, Timothy A.
 APPLICANT: Tumas, Daniel
 APPLICANT: Watanabe, Colin K
 APPLICANT: Wood, William
 APPLICANT: Zhang
 TITLE OF INVENTION:
 FILE REFERENCE:
 CURRENT APPLICATION NUMBER: US/10/028,072
 CURRENT FILING DATE: 2001-12-19
 PRIOR APPLICATION NUMBER: 60/049911
 PRIOR FILING DATE: 1997-06-18
 PRIOR APPLICATION NUMBER: 60/056974
 PRIOR FILING DATE: 1997-08-26
 PRIOR APPLICATION NUMBER: 60/059113
 PRIOR FILING DATE: 1997-09-17
 PRIOR APPLICATION NUMBER: 60/059115
 PRIOR FILING DATE: 1997-09-17
 PRIOR APPLICATION NUMBER: 60/059117
 PRIOR FILING DATE: 1997-09-17
 PRIOR APPLICATION NUMBER: 60/059122
 PRIOR FILING DATE: 1997-09-17
 PRIOR APPLICATION NUMBER: 60/059184
 PRIOR FILING DATE: 1997-09-17
 PRIOR APPLICATION NUMBER: 60/059263
 PRIOR FILING DATE: 1997-09-18
 PRIOR APPLICATION NUMBER: 60/059352
 PRIOR FILING DATE: 1997-09-19
 PRIOR APPLICATION NUMBER: 60/059588
 PRIOR FILING DATE: 1997-09-19
 PRIOR APPLICATION NUMBER: 60/059836
 PRIOR FILING DATE: 1997-09-24
 PRIOR APPLICATION NUMBER: 60/062250
 PRIOR FILING DATE: 1997-10-17
 PRIOR APPLICATION NUMBER: 60/062285
 PRIOR FILING DATE: 1997-10-17
 PRIOR APPLICATION NUMBER: 60/062287
 PRIOR FILING DATE: 1997-10-17
 PRIOR APPLICATION NUMBER: 60/062814
 PRIOR FILING DATE: 1997-10-24
 PRIOR APPLICATION NUMBER: 60/062816
 PRIOR FILING DATE: 1997-10-24
 PRIOR APPLICATION NUMBER: 60/063045
 PRIOR FILING DATE: 1997-10-24
 PRIOR APPLICATION NUMBER: 60/063082
 PRIOR FILING DATE: 1997-10-31
 PRIOR APPLICATION NUMBER: 60/063127
 PRIOR FILING DATE: 1997-10-24
 PRIOR APPLICATION NUMBER: 60/063327
 PRIOR FILING DATE: 1997-10-27
 PRIOR APPLICATION NUMBER: 60/063329
 PRIOR FILING DATE: 1997-10-27
 PRIOR APPLICATION NUMBER: 60/063550
 PRIOR FILING DATE: 1997-10-28
 PRIOR APPLICATION NUMBER: 60/063561
 PRIOR FILING DATE: 1997-10-28
 PRIOR APPLICATION NUMBER: 60/063704
 PRIOR FILING DATE: 1997-10-29
 PRIOR APPLICATION NUMBER: 60/063733
 PRIOR FILING DATE: 1997-10-29
 PRIOR APPLICATION NUMBER: 60/063735
 PRIOR FILING DATE: 1997-10-29
 PRIOR APPLICATION NUMBER: 60/063738
 PRIOR FILING DATE: 1997-10-29
 PRIOR APPLICATION NUMBER: 60/063755
 PRIOR FILING DATE: 1997-10-17
 PRIOR APPLICATION NUMBER: 60/064248
 PRIOR FILING DATE: 1997-11-03
 PRIOR APPLICATION NUMBER: 60/064809
 PRIOR FILING DATE: 1997-11-07
 PRIOR APPLICATION NUMBER: 60/065186
 PRIOR FILING DATE: 1997-11-12
 PRIOR APPLICATION NUMBER: 60/065846
 PRIOR FILING DATE: 1997-11-17

PRIOR APPLICATION NUMBER: 60/066364
 PRIOR FILING DATE: 1997-11-21
 PRIOR APPLICATION NUMBER: 60/066453
 PRIOR FILING DATE: 1997-11-24
 PRIOR APPLICATION NUMBER: 60/066511
 PRIOR FILING DATE: 1997-11-24
 PRIOR APPLICATION NUMBER: 60/066770
 PRIOR FILING DATE: 1997-11-24
 PRIOR APPLICATION NUMBER: 60/069212
 PRIOR FILING DATE: 1997-12-11
 PRIOR APPLICATION NUMBER: 60/069278
 PRIOR FILING DATE: 1997-12-11
 PRIOR APPLICATION NUMBER: 60/069334
 PRIOR FILING DATE: 1997-12-11
 PRIOR APPLICATION NUMBER: 60/069694
 PRIOR FILING DATE: 1997-12-16
 PRIOR APPLICATION NUMBER: 60/072320
 PRIOR FILING DATE: 1998-01-23
 PRIOR APPLICATION NUMBER: 60/073612
 PRIOR FILING DATE: 1998-02-04
 PRIOR APPLICATION NUMBER: 60/074086
 PRIOR FILING DATE: 1998-02-09
 PRIOR APPLICATION NUMBER: 60/074092
 PRIOR FILING DATE: 1998-02-09
 PRIOR APPLICATION NUMBER: 60/077791
 PRIOR FILING DATE: 1998-03-12
 PRIOR APPLICATION NUMBER: 60/078910
 PRIOR FILING DATE: 1998-03-20
 PRIOR APPLICATION NUMBER: 60/079294
 PRIOR FILING DATE: 1998-03-25
 PRIOR APPLICATION NUMBER: 60/079663
 PRIOR FILING DATE: 1998-02-27
 PRIOR APPLICATION NUMBER: 60/079728
 PRIOR FILING DATE: 1998-03-27
 PRIOR APPLICATION NUMBER: 60/080165
 PRIOR FILING DATE: 1998-03-31
 PRIOR APPLICATION NUMBER: 60/081203
 PRIOR FILING DATE: 1998-04-09
 PRIOR APPLICATION NUMBER: 60/081229
 PRIOR FILING DATE: 1998-04-09
 PRIOR APPLICATION NUMBER: 60/081695
 PRIOR FILING DATE: 1998-04-14
 PRIOR APPLICATION NUMBER: 60/081817
 PRIOR FILING DATE: 1998-04-15
 PRIOR APPLICATION NUMBER: 60/081818
 PRIOR FILING DATE: 1998-04-15
 PRIOR APPLICATION NUMBER: 60/082999
 PRIOR FILING DATE: 1998-04-24
 PRIOR APPLICATION NUMBER: 60/083322
 PRIOR FILING DATE: 1998-04-28
 PRIOR APPLICATION NUMBER: 60/083545
 PRIOR FILING DATE: 1998-04-29
 PRIOR APPLICATION NUMBER: 60/084600
 PRIOR FILING DATE: 1998-05-07
 PRIOR APPLICATION NUMBER: 60/084627
 PRIOR FILING DATE: 1998-05-07
 PRIOR APPLICATION NUMBER: 60/084637
 PRIOR FILING DATE: 1998-05-07
 PRIOR APPLICATION NUMBER: 60/085149
 PRIOR FILING DATE: 1998-05-12
 PRIOR APPLICATION NUMBER: 60/085323
 PRIOR FILING DATE: 1998-05-13
 PRIOR APPLICATION NUMBER: 60/085338
 PRIOR FILING DATE: 1998-05-13
 PRIOR APPLICATION NUMBER: 60/085339
 PRIOR FILING DATE: 1998-05-13
 PRIOR APPLICATION NUMBER: 60/085579
 PRIOR FILING DATE: 1998-05-15
 PRIOR APPLICATION NUMBER: 60/085697
 PRIOR FILING DATE: 1998-05-15
 PRIOR APPLICATION NUMBER: 60/085704
 PRIOR FILING DATE: 1998-05-15
 PRIOR APPLICATION NUMBER: 60/086414

```
; PRIOR FILING DATE: 1998-05-22
; PRIOR APPLICATION NUMBER: 60/086430
; PRIOR FILING DATE: 1998-05-22
; PRIOR APPLICATION NUMBER: 60/087106
; PRIOR FILING DATE: 1998-05-28
; PRIOR APPLICATION NUMBER: 60/088026
; PRIOR FILING DATE: 1998-06-04
; PRIOR APPLICATION NUMBER: 60/088730
; PRIOR FILING DATE: 1998-06-10
; PRIOR APPLICATION NUMBER: 60/088741
; PRIOR FILING DATE: 1998-06-10
; PRIOR APPLICATION NUMBER: 60/088810
; PRIOR FILING DATE: 1998-06-10
; PRIOR APPLICATION NUMBER: 60/088858
; PRIOR FILING DATE: 1998-06-11
; PRIOR APPLICATION NUMBER: 60/089332
; PRIOR FILING DATE: 1998-06-17
; PRIOR APPLICATION NUMBER: 60/089599
; PRIOR FILING DATE: 1998-06-17
; PRIOR APPLICATION NUMBER: 60/089907
; PRIOR FILING DATE: 1998-06-18
; PRIOR APPLICATION NUMBER: 60/089947
; PRIOR FILING DATE: 1998-06-19
; PRIOR APPLICATION NUMBER: 60/090349
; PRIOR FILING DATE: 1998-06-23
; PRIOR APPLICATION NUMBER: 60/090429
; PRIOR FILING DATE: 1998-06-24
; PRIOR APPLICATION NUMBER: 60/090445
; PRIOR FILING DATE: 1998-06-24
; PRIOR APPLICATION NUMBER: 60/090538
; PRIOR FILING DATE: 1998-06-24
; PRIOR APPLICATION NUMBER: 60/090863
; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: 60/091360
; PRIOR FILING DATE: 1998-07-01
; PRIOR APPLICATION NUMBER: 60/091519
; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
```

Query Match 97.7%; Score 1949; DB 9; Length 364;

Best Local Similarity 98.4%; Pred. No. 1.9e-157; Indels 6; Gaps 1;

Matches 364; Conservative 0; Mismatches 0; Indels 6; Gaps 1;

```
QY 1 MHRLLFYVTLICANFCSCRDTSATPQASISIKALRNANLRDESNHLTDLRRDETIOYKG 60
DB 1 MHRLLFYVTLICANFCSCRDTSATPQASISIKALRNANLRD-----DLYRDETIOYKG 54
QY 61 NGYVSPRPFPNSYPNNLLTWRLHSQENTRIQLVFDNOFGLBEAENDICRDVEVEDIS 120
DB 55 NGYVSPRPFPNSYPNNLLTWRLHSQENTRIQLVFDNOFGLBEAENDICRDVEVEDIS 114
QY 121 ESTTIIRGRCGKHEVPPRIKSRITNQIKITFKSDDYFYAKPGFKIYYSILEDFOPAASE 180
DB 115 ESTTIIRGRCGKHEVPPRIKSRITNQIKITFKSDDYFYAKPGFKIYYSILEDFOPAASE 174
QY 181 TNMESVTSISIGSVSNPSVTDPTLIADALDKKIAEFTVEDLKYFNPESMOEDLENNY 240
DB 175 TNMESVTSISIGSVSNPSVTDPTLIADALDKKIAEFTVEDLKYFNPESMOEDLENNY 234
QY 241 LDTFRYRGRSYHDRKSKYDLDRLNDADARYSCTPNYSVNI REELKLANVFFPCCLLYO 300
DB 235 LDTFRYRGRSYHDRKSKYDLDRLNDADARYSCTPNYSVNI REELKLANVFFPCCLLYO 294
QY 301 RCGNCGCGTVMNRSCNCGTVMNRSCNCKTYKYEVLQFEFGHKKRRGRATMALVLDIQLDHHRC 360
DB 295 RCGNCGCGTVMNRSCNCGTVMNRSCNCKTYKYEVLQFEFGHKKRRGRATMALVLDIQLDHHRC 354
QY 361 DCICSSRPPR 370
DB 355 DCICSSRPPR 364
```

RESULT 12

US-10-121-049-186

; Sequence 186, Application US/10121049

; Publication No. US2003002239A1

; GENERAL INFORMATION:

; APPLICANT: Baker, Kevin P.

; APPLICANT: Beresini, Maureen

; APPLICANT: Deforge, Laura

; APPLICANT: Desnoyers, Luc

; APPLICANT: Filvaroff, Ellen

; APPLICANT: Gao, Wei-Qiang

; APPLICANT: Gerritsen, Mary E.

; APPLICANT: Goddard, Audrey

; APPLICANT: Godowski, Paul J.

; APPLICANT: Gurney, Austin L.

; APPLICANT: Sherwood, Steven

; APPLICANT: Smith, Victoria

; APPLICANT: Stewart, Timothy A.

; APPLICANT: Tamas, Daniel

; APPLICANT: Watanabe, Colin K

; APPLICANT: Wood, William

; APPLICANT: Zhang, Zemin

; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC

; TITLE OF INVENTION: ACIDS ENCODING THE SAME

; FILE REFERENCE: P3330R1C17

; CURRENT FILING DATE: 2002-04-12

; PRIOR APPLICATION NUMBER: US/10/121,049

; PRIOR APPLICATION REMOVED - See File Wrapper or Palm

; NUMBER OF SEQ ID NOS: 550

; SEQ ID NO 186

; LENGTH: 364

; TYPE: PRT

; ORGANISM: Homo Sapien

; US-10-121-049-186

Query Match 97.7%; Score 1949; DB 9; Length 364;

Best Local Similarity 98.4%; Pred. No. 1.9e-157; Indels 6; Gaps 1;

Matches 364; Conservative 0; Mismatches 0; Indels 6; Gaps 1;

```
QY 1 MHRLLFYVTLICANFCSCRDTSATPQASISIKALRNANLRDESNHLTDLRRDETIOYKG 60
DB 1 MHRLLFYVTLICANFCSCRDTSATPQASISIKALRNANLRD-----DLYRDETIOYKG 54
QY 61 NGYVSPRPFPNSYPNNLLTWRLHSQENTRIQLVFDNOFGLBEAENDICRDVEVEDIS 120
DB 55 NGYVSPRPFPNSYPNNLLTWRLHSQENTRIQLVFDNOFGLBEAENDICRDVEVEDIS 114
QY 121 ESTTIIRGRCGKHEVPPRIKSRITNQIKITFKSDDYFYAKPGFKIYYSILEDFOPAASE 180
DB 115 ESTTIIRGRCGKHEVPPRIKSRITNQIKITFKSDDYFYAKPGFKIYYSILEDFOPAASE 174
QY 181 TNMESVTSISIGSVSNPSVTDPTLIADALDKKIAEFTVEDLKYFNPESMOEDLENNY 240
DB 175 TNMESVTSISIGSVSNPSVTDPTLIADALDKKIAEFTVEDLKYFNPESMOEDLENNY 234
QY 241 LDTFRYRGRSYHDRKSKYDLDRLNDADARYSCTPNYSVNI REELKLANVFFPCCLLYO 300
DB 235 LDTFRYRGRSYHDRKSKYDLDRLNDADARYSCTPNYSVNI REELKLANVFFPCCLLYO 294
QY 301 RCGNCGCGTVMNRSCNCGTVMNRSCNCKTYKYEVLQFEFGHKKRRGRATMALVLDIQLDHHRC 360
DB 295 RCGNCGCGTVMNRSCNCGTVMNRSCNCKTYKYEVLQFEFGHKKRRGRATMALVLDIQLDHHRC 354
QY 361 DCICSSRPPR 370
DB 355 DCICSSRPPR 364
```

RESULT 13

US-10-123-904-186

; Sequence 186, Application US/10123904

; Publication No. US20030022328A1

; GENERAL INFORMATION:

```

; APPLICANT: Baker, Kevin P.
; APPLICANT: Beresini, Maureen
; APPLICANT: Deforge, Laura
; APPLICANT: Desnoyers, Luc
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Sherwood, Steven
; APPLICANT: Smith, Victoria
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K
; APPLICANT: Wood, William
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; FILE REFERENCE: P330R1C54
; CURRENT APPLICATION NUMBER: US/10/123,904
; PRIOR APPLICATION REMOVED - See File Wrapper or Palm
; NUMBER OF SEQ ID NOS: 550
; SEQ ID NO 186
; LENGTH: 364
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-123-904-186

```

```

Query Match          97.7%; Score 1949; DB 9; Length 364;
Best Local Similarity 98.4%; Pred. No. 1.9e-157;
Matches 364; Conservative 0; Mismatches 0; Indels 6; Gaps 1;

```

```

QY 1 MHRLEFYTLICANFCSCRDTSATPOSASIKALRNANLRDESNHLLTDLYRDETTQVKG 60
DB 1 MHRLEFYTLICANFCSCRDTSATPOSASIKALRNANLRD-----DLRRDETTQVKG 54
QY 61 NGVOSRPFNSYPRLNLLTWRLHSEENTRIQLVFNQFLEAENDICRYDEVEVDIS 120
DB 55 NGVOSRPFNSYPRLNLLTWRLHSEENTRIQLVFNQFLEAENDICRYDEVEVDIS 114
QY 121 ETSITIRGRMGCKHEVPPRIKSRNTQIKITFKSDDYFVANPGRKIYSLLEDPQPAASE 180
DB 115 ETSITIRGRMGCKHEVPPRIKSRNTQIKITFKSDDYFVANPGRKIYSLLEDPQPAASE 174
QY 181 TMSVSTSSISGVSYNSPSVTDPDLLADALDKKIAEDYEDLLKTFNPESMOEDLENMY 240
DB 175 TMSVSTSSISGVSYNSPSVTDPDLLADALDKKIAEDYEDLLKTFNPESMOEDLENMY 234
QY 241 LDPFRYRGRSYHDKRSKVDLDRLNDADAKRYSCTPRNYSVIRBELKIANVFFPCLLYQ 300
DB 235 LDPFRYRGRSYHDKRSKVDLDRLNDADAKRYSCTPRNYSVIRBELKIANVFFPCLLYQ 294
QY 301 RCGNGCGGTVMNRSCCTNSGKTVKKYHEVLOFEPGHIKRGRAKTMALVDIOLDHNERC 360
DB 295 RCGNGCGGTVMNRSCCTNSGKTVKKYHEVLOFEPGHIKRGRAKTMALVDIOLDHNERC 354
QY 361 DCICSSRPPR 370
DB 355 DCICSSRPPR 364

```

```

RESULT 14
US-10-140-470-186
; Sequence 186, Application US/10140470
; Publication No. US20030022331A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
; APPLICANT: Beresini, Maureen
; APPLICANT: Deforge, Laura
; APPLICANT: Desnoyers, Luc
; APPLICANT: Filvaroff, Ellen

```

```

; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Sherwood, Steven
; APPLICANT: Smith, Victoria
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K
; APPLICANT: Wood, William
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; FILE REFERENCE: P330R1C160
; CURRENT APPLICATION NUMBER: US/10/140,470
; PRIOR APPLICATION REMOVED - See Palm or File Wrapper
; NUMBER OF SEQ ID NOS: 550
; SEQ ID NO 186
; LENGTH: 364
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-140-470-186

```

```

Query Match          97.7%; Score 1949; DB 9; Length 364;
Best Local Similarity 98.4%; Pred. No. 1.9e-157;
Matches 364; Conservative 0; Mismatches 0; Indels 6; Gaps 1;

```

```

QY 1 MHRLEFYTLICANFCSCRDTSATPOSASIKALRNANLRDESNHLLTDLYRDETTQVKG 60
DB 1 MHRLEFYTLICANFCSCRDTSATPOSASIKALRNANLRD-----DLRRDETTQVKG 54
QY 61 NGVOSRPFNSYPRLNLLTWRLHSEENTRIQLVFNQFLEAENDICRYDEVEVDIS 120
DB 55 NGVOSRPFNSYPRLNLLTWRLHSEENTRIQLVFNQFLEAENDICRYDEVEVDIS 114
QY 121 ETSITIRGRMGCKHEVPPRIKSRNTQIKITFKSDDYFVANPGRKIYSLLEDPQPAASE 180
DB 115 ETSITIRGRMGCKHEVPPRIKSRNTQIKITFKSDDYFVANPGRKIYSLLEDPQPAASE 174
QY 181 TMSVSTSSISGVSYNSPSVTDPDLLADALDKKIAEDYEDLLKTFNPESMOEDLENMY 240
DB 175 TMSVSTSSISGVSYNSPSVTDPDLLADALDKKIAEDYEDLLKTFNPESMOEDLENMY 234
QY 241 LDPFRYRGRSYHDKRSKVDLDRLNDADAKRYSCTPRNYSVIRBELKIANVFFPCLLYQ 300
DB 235 LDPFRYRGRSYHDKRSKVDLDRLNDADAKRYSCTPRNYSVIRBELKIANVFFPCLLYQ 294
QY 301 RCGNGCGGTVMNRSCCTNSGKTVKKYHEVLOFEPGHIKRGRAKTMALVDIOLDHNERC 360
DB 295 RCGNGCGGTVMNRSCCTNSGKTVKKYHEVLOFEPGHIKRGRAKTMALVDIOLDHNERC 354
QY 361 DCICSSRPPR 370
DB 355 DCICSSRPPR 364

```

```

RESULT 15
US-10-175-746-186
; Sequence 186, Application US/10175746
; Publication No. US20030027270A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
; APPLICANT: Beresini, Maureen
; APPLICANT: Deforge, Laura
; APPLICANT: Desnoyers, Luc
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Gurney, Austin L.

```

```

: APPLICANT: Sherwood, Steven
: APPLICANT: Smith, Victoria
: APPLICANT: Stewart, Timothy A.
: APPLICANT: Tumas, Daniel
: APPLICANT: Watanabe, Colin K
: APPLICANT: Wood, William
: APPLICANT: Zhang, Zemin
: TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
: TITLE OF INVENTION: ACIDS ENCODING THE SAME
: FILE REFERENCE: P330R1C353
: CURRENT APPLICATION NUMBER: US/10/175,746
: CURRENT FILING DATE: 2002-06-19
: Prior Application removed - See File Wrapper or Palm
: NUMBER OF SEQ ID NOS: 550
: SEQ ID NO 186
: LENGTH: 364
: TYPE: PRT
: ORGANISM: Homo Sapien
US-10-175-746-186

```

Query Match 97.7%; Score 1949; DB 9; Length 364;

Best Local Similarity 98.4%; Pred. No. 1.9e-157; Mismatches 0; Indels 6; Gaps 1;

```

Db 1 MHRLLFYTLICANFCSCROTSATPOSASIKALRNANLRDESNHLTDLYRDETIQYKG 60
    1 MHRLLFYTLICANFCSCROTSATPOSASIKALRNANLRD-----DLYRDETIQYKG 54
QY 61 NGYVQSPREFPNSTYRNLLTWRLHSQENTRIQLVFDNQGLEAENDICRYDFVEVEDIS 120
    55 NGYVQSPREFPNSTYRNLLTWRLHSQENTRIQLVFDNQGLEAENDICRYDFVEVEDIS 114
Db 121 ETSITLGRMGCHKEVPRIRKSTNQIKITFKSDDYFAKPGFKIYSLLEDFOPAASE 180
    115 ETSITLGRMGCHKEVPRIRKSTNQIKITFKSDDYFAKPGFKIYSLLEDFOPAASE 174
QY 181 TNMESVTSISIGSVNSPSVTDPTLIADALDKIAEFTVEDLKYFNPESMOEDLENNY 240
    175 TNMESVTSISIGSVNSPSVTDPTLIADALDKIAEFTVEDLKYFNPESMOEDLENNY 234
Db 241 LDTPRYRGRSYHDKRSKYDLRLNDDAKRYSCTPRNSVNIREEIKLANVFFPRCLLVQ 300
    235 LDTPRYRGRSYHDKRSKYDLRLNDDAKRYSCTPRNSVNIREEIKLANVFFPRCLLVQ 294
QY 301 RCGNCGCGTNNMRSCTGNSGKTYKKYHEVLQEPFGHKKRGRAKTMALVDIQLDHHERC 360
    295 RCGNCGCGTNNMRSCTGNSGKTYKKYHEVLQEPFGHKKRGRAKTMALVDIQLDHHERC 354
Db 361 DCICSSRPPR 370
    355 DCICSSRPPR 364

```

Search completed: June 12, 2003, 15:40:45
Job time : 25 secs

